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|   | equence 6,   | equence 8,                            | equence 9,<br>equence 10.               | equence 11,                               | equence 457   | equence 4/35,                           | equence 4682                             | equence 14346,                           | equence 5                                | equence 751                              | equence 577                             | equence 5,                               | equenc                                 | equence 563                               | equence 15,                             | equence 29,                              | equence 9,                              | equence 10,                              | equence 11,                              | equence 30,                              | equence 3.                               | odenie 2                                | י<br>טור<br>טור  | equence 6,   | equence 7,                               | א פיזורפווה פ                            | ים המתחומים                               | equence 9,  | equence 10,                            | equence 427                           | Ant annatha                          |   | equence 4569,  | se 592                                | edience 6203                            | equence ozon,                             | ore annanha  | equenc                                  | equence 31,                              | equence 6733                             | A annama                                 | 117 00000000000000000000000000000000000  | יהלתפוונים להי              | '*cd aouanba   | Φ  | equence 2400                         | equence 372,                          | e 2238                                     | equence 3869                       | a charge                              | 000000000000000000000000000000000000000  | eduence 23                | ri acuanha   | equence 18  | equence 60                            | e 54                                     | equence 48                            | equence 47                               | equence 99                            | אסטופינהם                             | בלתכווכם בי                             | 'c aouanba                           | equence 5,                            | equence 5,                                | equence 5,                             | equence 5                               | Panence 67                                 |   | equence 67                                  | eguence 49                                  | equence 67                             | Cy appearing                           | ence 67,                               | equence 5298                           | Act opiois                              |
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| Sequence 92,<br>Sequence 92,<br>Patent No. 5;<br>Sequence 14,   | Sequence 14,   | Sequence 1,                           | Sequence 57                             | 3 Sequence 57                             | Sequence 35   | Sequence 35                             | Sequence 59                              | Sequence 128                             | Sequence 598                             | Seguence 37,                             | Sequence 35                             | Sequence 37,                             | Seguence 1,                            | Sequence 44.                              | Sednence 19,                            | Seguence 178                             | Sequence 181                            | Sequence 38,                             | Seguence 38,                             | Sequence 53,                             | Sequence 54,                             | Semenas 57                              | ' C aniinhag   | sednence 23'   | Sequence 54,                             | Seguence 57                              | Section 15                                | 'cī anmanhac  | Sequence 1,                            | Sequence 1,                           | Seminary 1                           | ייי סטונטנטטטטטטטטטטטטטטטטטטטטטטטטטטטטטטטטט                         | 'I ednence I'  | Sequence 1,                           | Semenas 16                              | or company                                | 'AT DOMESTIC TO  | sednence 18'                            | Sequence 19,                             | Sequence 20,                             | Segmence 21                              | Security Sec | CC obtoined                 | seducine 23  | Sequence II'                             | 's edneuce 3'                        | Seguence 8,                           | 96 Seguence 600                            | Patent No. 51                      | Semience 1.                           | Company 1                                | Sequence 1,               | מפלתפווכם די   | 'T eagnement'   | Sednence 2,                           | Sequence 33,                             | Sequence 1,                           | Sequence 153                             | Sequence 2,                           | Semence 2.                            | C CONTONION                             | Seducince 2,                         | sednence z,                           | Sequence 408                              | Sequence 2,                            | Seguence 46.                            | Seguence 4619                              | THOS COMORDO  | sednence 4/65,                              | Sequence 38, Ap                             | Sequence 6,                            | Semience 7                             | '/ ealleance                           | Sequence 8,                            | Compress of                             |
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| 2 US-10-676-475A-92 Sequence 92, 2 US-10-607-595-92 Sequence 92, 6 5304631-9 Patent No. 55, 1 US-08-400-796-14 Sequence 14, 1 US-08-400-796-14          | I US-U6-248-839C-144 Sequence 14, 2 US-09-567-003C-4 Sequence 4, | 1 US-08-846-526-1 Sequence 1,         | 1 US-08-800-644-57 Sequence 57          | 2 US-09-438-185A-578 Sequence 570         | 2 US-09-911-92/-35 Sequence 35,                                 | 2 US-09-911-888-35 Sequence 35          | 2 US-09-205-258-598 Sequence 59          | 2 US-09-079-030-128 Sequence 128         | 2 US-10-004-860-598 Sequence 598         | 1 US-08-690-011A-37 Sequence 37,         | 2 US-08-965-762-35 Sequence 35,         | 2 US-09-299-495F-37 Sequence 37          | 1 US-08-549-752A-1 Sequence 1,         | 2 US-U6-838-20/A-442 Sequence 445         | 2 US-09-756-983-19 Sequence 19,         | 2 US-09-9/3-2/8-1/8 Sequence 178         | 2 US-09-227-357-181 Sequence 18         | 1 US-08-690-011A-38 Sequence 38,         | 2 US-09-299-495F-38 Sequence 38,         | 1 US-08-690-011A-53 Sequence 53,         | 1 US-08-690-011A-54 Sequence 54,         | 1 US-08-690-011A-57 Semience 57         | 2 118-00-300-400B-B3   | 2 US-US-239-492F-33 Sequence 33,   | 2 US-09-299-495F-54 Sequence 54,         | 2 US-09-299-495F-57 Sequence 57          | 2 TIS-09-839-5773-15                      | t to or or and the To   | 1 US-08-491-527A-1 Sequence 1,         | 1 US-08-845-659-1 Seguence 1,         | 2 US-08-845-658-1 Semience 1         | 2 TG 00 E10 170 1   | 'T earners T-8-T-8-C-61'   | 2 US-09-517-597-1 Sequence 1,         | 2 US-08-751-344B-16 Semience 16         | 2 TRE-08-751-2440-17                      | 2 10 00 101 01 01 01 01 00 00  | 2 02-00-121-3445-16 Sequence 18,        | 7 OS-08-721-344B-19 Sequence 19,         | 2 US-08-751-344B-20 Sequence 20,         | 2 US-08-751-344B-21 Semience 21          | 2 IIS-08-751-1448-22   | 2 TRC_08_751_244B_23        | מביס שפקעפווכפ ליי ביי ייי פפקעפווכפ ליי ביי ייי ייי ייי ייי יייי יייי יייי  | 2 US-U9-395-UI/B-II Sequence II,         | 2 US-U9-39U-85I-3 Sequence 3,        | 2 US-09-390-851-8 Sequence 8,         | 2 US-09-270-767-60096 Seguence 600         | 6 5177197-34 Patent No. 51         | 2 US-08-393-889-1 Semience 1.         | 2 TR-00-110-000-1                        | 2 UB-03-113-200-1         | 1 00-00-3/E-00-00  | 4 FCI-0394-09450-I  | 2 US-U9-921-144-2                     | 2 US-08-960-190A-33 Sequence 33,         | 1 US-07-835-860-1 Sequence 1,         | 2 US-09-314-268-153 Sequence 153         | 1 US-07-940-861-2 Sequence 2,         | 1 US-08-459-512-2 Segmence 2.         | 1 110-00-4E0-4E0-6E7-0                  | 1 00-00-403-03/-2                    | 1 US-US-46U-132-2                     | Z US-09-621-976-4080 Sequence 408         | 4 PCT-US92-02050-2 Sequence 2,         | 2 US-09-277-078-46 Seguence 46.         | 2 US-09-621-976-4619 Segmence 4619         | 0 10 - 00 - 01 - 01 - 01 - 01 - 01 - 01               | 2 US-US-621-9/6-4/65 Sequence 4/65,         | 1 US-08-480-473B-38 Sequence 38, Ap         | 1 US-08-549-004A-6 Sequence 6,         | 1 US-08-549-004A-7 Semience 7          | 1 02-00-549-004A-7 Sequence 7,         | 1 US-08-549-004A-8 Sequence 8,         | 1 115-08-549-004b-9                     |
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| ក្នុងបង្គុស្ត្រប្រជុំក្នុងក្នុងមួយប្រជុំក្នុងប្រជុំក្នុងប្រជុំក្នុងក្នុងក្នុងក្នុងក្នុងក្នុងក្នុងប្រជុំក្នុងប្រជុំក្នុងប្រជុំក្នុងប្រជុំក្នុងប្រជុំក្នុងប្រជុំក្នុងប្រជុំក្នុងប្រជុំក្នុងប្រជុំក្នុងប្រជុំក្នុងប្រជុំក្នុងប្រជុំក្នុងប្រជុំក្នុងប្រជុំក្នុងប្រភពិធីក្នុងប្រជុំក្រុងប្រជុំក្នុងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រស្នងប្រជុំក្រុងប្រងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រុងប្រជុងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រុងប្រជុំក្រងប្រជុំក្រាងប្រជុំក្រងប្រជុំក្រងប្រជុំក្រងប្រជុំក្រងប្រជុងប្រជុងប្រជុងប្រជុំក្រងប្រជុំក្រងប្បក្សប្រជុំក្រងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុំក្រងប្រជុងប្រជុំក្រងប្បក្សប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្រជុងប្ប | 824, A<br>5821,<br>Appl<br>1662,<br>Appl<br>3, App<br>3, Appl<br>Appl<br>Appl<br>Appl<br>Appl<br>Appl<br>Appl   |
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| US-09-071-035-10 US-09-134-000C-6 US-10-206-576-10 US-09-095-117-2 US-09-095-117-4 US-09-095-117-4 US-09-095-117-4 US-09-095-117-4 US-09-095-117-4 US-09-095-117-4 US-09-095-117-4 US-09-905-107-532A-6 US-09-905-2-991A-2 US-09-905-2-991A-2 US-09-905-2-991A-2 US-09-905-2-991A-3 US-09-905-2-991A-3 US-09-905-2-991A-3 US-09-905-2-991A-3 US-09-905-2-991A-3 US-09-905-2-991A-3 US-09-905-2-991A-3 US-09-905-2-991A-3 US-09-252-991A-3 US-09-252-991A-3 US-09-252-991A-3 US-09-252-991A-3 US-09-252-991A-3 US-09-252-991A-3 US-09-252-991A-3 US-09-252-991A-3 US-09-252-991A-3 US-09-270-767-41   | 0 - SU<br>0 |
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| US-09-252-991A-22716 US-09-674-529B-14 US-09-544-529B-14 US-09-248-796A-1568 US-09-949-016-9718 US-09-949-016-9718 US-09-949-016-9718 US-09-949-016-9718 US-09-949-016-9718 US-09-949-016-9719 US-09-949-016-9719 US-09-134-001C-4400 US-09-949-016-9719 US-09-949-016-0451 US-09-949-016-0523   | US-09-248-796A-19891<br>US-09-538-092-352<br>US-09-298-894-29<br>US-09-734-237B-50<br>US-09-734-237B-50<br>US-09-734-237B-50<br>US-09-734-237B-51<br>US-09-232-197-51<br>US-09-232-197-51<br>US-09-232-197-51<br>US-09-232-197-51<br>US-09-232-197-51<br>US-09-232-197-51<br>US-09-232-197-51<br>US-09-232-197-51<br>US-09-232-198-51<br>US-09-232-198-51<br>US-09-562-991A-28389<br>US-09-562-737-6  |
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; ORGANISM: Streptococcus equi
US-09-103-664A-2
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Best Local Similarity 100.
Matches 9; Conservative
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296 RAAFLEKEN 304
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TYPE: PRT
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; Sequence 10895, Application US/09949016
; Batent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: CLOOD1307
; CURRENT APPLICATION NUMBER: US/09/949,016
; CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-03
; PRIOR FILING DATE: 2000-10-03
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FREESEQ for Windows Version 4.0
; SEQ ID NO 10895
; LENGTH: 345
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APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
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                                Sequence 251, App
Sequence 43, Appl
Sequence 2745, App
Sequence 197, App
Sequence 37129, A
Sequence 37129, A
Sequence 3794, App
Sequence 197, App
Sequence 197, App
Sequence 8, Appli
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8, Appli
3174, Ap
17293, A
12, Appl
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US-09-949-016-11550

US-09-878-574-2

US-09-326-43

US-09-252-991A-27426

US-09-148-545-197

US-09-148-545-197

US-09-270-76-53146

US-09-270-76-53346

US-09-107-433-3794

US-09-52-991A-2828

US-09-52-991A-2828

US-09-999-016-1136

US-09-999-016-1136

US-09-999-016-1136

US-09-377-2858-12

US-09-377-2858-12

US-09-377-2858-12

US-09-377-2858-12

US-09-377-2858-12
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; Sequence 11210, Application US/09949016
; Patent No. 6812339
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Best Local Similarity 100.
Matches 19; Conservative
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ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-949-016-10895
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Sequence 2, Application US/09103664A

Patent No. 6458358
GENERAL INFORMATION:
APPLICANT: University of Kentucky Research Foundation
APPLICANT: Timoney, John
APPLICANT: Artiushin, Sergey
TITLE OF INVENTION: and Assays Therefor
FILE REFERENCE: 50229-212
CURRENT FILLIG DATE: 1998-06-23
CURRENT FILLIG DATE: 1998-06-23
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin version 3.0
SOFTWARE: Patentin version 3.0
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;Patent No. 5290690
;Patent No. 5290690
;MATHYSENS, IGNACE;STANSSENS, PATRICK
;MATHYSENS, GASTON WODAK, SHOSHANA, QUAX, WILHELMUS J.
;TITLE OF INVENTION: METHODS AND MEANS FOR CONTROLLING THE
;STABILITY OF PROTEINS
; NUMBER OF SEQUENCES: 22
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/398,706
; FILING DATE: 25-AUG-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 334;
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100.0%; Pred. No. 0.086;
Live 0; Mismatches 0
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100.0%; Pred. No. 1;
tive 0; Mismatches
CURRENT FILING DATE: 2000-04-14

PRIOR APPLICATION NUMBER: 6/241,755

PRIOR FILING DATE: 2000-10-20

PRIOR PELICATION NUMBER: 60/237,768

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-0-03

PRIOR FILING DATE: 2000-09-08

NUMBER OF SEQ ID NOS: 207012

SOFTWARE: FastSEQ for Windows Version 4.0

LENGTH: 335
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Sequence 5144, Application US/09583110
Sequence 5144, Application US/09583110
Patent No. 6699703
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Nucleic Acid and Amino Acid Sequences Relating to Streptococcus
TITLE OF INVENTION: Pretamoniae for Diagnostics and Therapeutics
TITLE OF INVENTION: Pretamoniae for Diagnostics and Therapeutics
TITLE OF INVENTION: Number: US/09/583,110
CURRENT APPLICATION NUMBER: US 99/107,433
PRIOR APPLICATION NUMBER: US 60/085,131
PRIOR FILING DATE: 1998-06-30
PRIOR FILING DATE: 1998-06-30
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Patent No. 6617156
GENERAL INFORMATION:
APPLICANT: Lynn Doucette-Stamm et al
APPLICANT: Lynn Doucette-Stamm et al
APPLICANT: Lynn Doucette-Stamm et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
TITLE OF INVENTION: NUMBER: 02/09/134,000C
CURRENT PILING DATE: 1998-08-13
PRIOR APPLICATION NUMBER: US 60/055,778
PRIOR APPLICATION NUMBER: US 60/055,778
PRIOR FILING DATE: 1997-08-15
NUMBER OF SEQ ID NOS: 6812
SOFTWARE: Patentin version 3.1
SEQ ID NO 5352
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Patent No. 63963

GENERAL INFORMATION:

APPLICANT: Dunas Milne Edwards, J.B.

APPLICANT: Uobert, S.

APPLICANT: Glordano, J.Y.

TITLE OF INVEXTION: ESTS and Encoded Human Proteins.

FILE REFERENCE: GENSET.054PR2.

CURRENT APPLICATION NUMBER: US/09/621,976

CURRENT APPLICATION NUMBER: US/09/621,976

CURRENT PILING DATE: 2000-07-21

NUMBER OF SEQ ID NOS: 19335

SEQ ID NO 4267

LENGTH: 95
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100.0%; Pred. No. 15;
tive 0; Mismatches
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100.0%; Pred. No. 38;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT;
CRGANISM: Enterococcus faecalis
US-09-134-000C-5352
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Best Local Similarity 100.
Matches 7; Conservative
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; ORGANISM: Homo sapiens
US-09-621-976-4267
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Sequence 45, Application US/09907907A

Sequence 45, Application US/09907907A

Sequence 45, Application US/09907907A

Sequence 45, Application

TITLE OF INVENTION: GENES DISPLAYING ENHANCED EXPRESSION DURING CELLULAR SENESCENCE A

TITLE OF INVENTION: TERMINAL CELL DIPFERRYIATION AND USES THEREOF

TITLE OF INVENTION: TERMINAL CELL DIPFERRYIATION AND USES THEREOF

FILE REPERENCE: A34584-A-PCT-USA (070050.1664)

CURRENT APPLICATION NUMBER: US/09/907,907A

PRIOR PILING DATE: 1999-02-02

NUMBER OF SEQ ID NOS: 51

SEQ ID NO 45

LENGTH: 245
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                                                                                                                                                                Sequence 16607, Application US/09902540

Patent No. 6833447

GENERAL INFORMATION:
APPLICANT: Goldman, Barry S.
APPLICANT: Misqand, Roger C.
APPLICANT: Slate, Steven C.
APPLICANT: MINGER: 1001549)B
CURRENT FILING DATE: 2000-07-10
PRIOR FILING DATE: 2000-07-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Consensus sequence, between Homo sapiens OLD-35 and ; OTHER INFORMATION: Bacillus subtilis PNPase US-09-907-907A-45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 84;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 24.1%; Score 7; DB 2; Best Local Similarity 100.0%; Pred. No. 8.4; Matches 7; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24.1%; Score 7; DB 2
100.0%; Pred. No. 2.9
ive 0; Mismatches
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US-09-134-000C-5352
; Sequence 5352, Application US/09134000C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Myxococcus xanthus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 16607
LENGTH: 84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.
Matches 7; Conservative
                                    391 VAELEKEV 398
       16 VAELEKEV 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       |||||||
70 RAAFLEK 76
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Length 112 0; Indels

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Sequence 5995, Application US/09134000C

Patent No. 6617156

GENERAL INFORMATION:

APPLICANT: Lynn Doucette-Stamm et al

APPLICANT: Lynn Doucette-Stamm et al

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES AND THERAPEUTICS

CURRENT APPLICATION NUMBER: US/09/134,000C

CURRENT FILING DATE: 1998-08-13

PRIOR FILING DATE: 1997-08-15

NUMBER OF SEQ ID NOS: 6812

SEQ ID NO 5995

INDIGHT: 121

MANNER NO 5995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: MISC_FEATURE

1. LOCATION: (35)

1. CHER INDEMATION: Amino acid 35 is Xaa wherein Xaa = any amino acid. US-09-134-000C-5995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Doxeey, Stephen J.
TITLE OF INVENTION: DIAGNOSTIC METHODS FOR SCREENING
TITLE OF INVENTION: PATIENTS FOR SCLERODERMA
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 225 Franklin Street
CITY: Boston
                                                                                           20.7%; Score 6; DB 2;
100.0%; Pred. No. 45;
ative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20.7%; Score 6; DB 2
100.0%; Pred. No. 48;
tive 0; Mismatches
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COUNTRY: US
ZIP: 02110-2804
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ VERSION 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/743,20
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 12, Application US/08743200; Patent No. 5861260; GENERAL INFORMATION:
      ; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-09-205-658-75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT ORGANISM: Enterococcus faecalis
                                                                       Query Match
Best Local Similarity 10v.v
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                                                                                                                                                                                                              28 AELEKE 33
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US-09-134-000C-5995
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                                                                                                                                                                                                                                                                                                                                                                                                                      Patent No. 6225120
GENERAL INFORMATION:
APPLICANT: Runkun, Gary
APPLICANT: Runkun, Gary
APPLICANT: Patterson, Gatth
APPLICANT: Posterson, Gatth
APPLICANT: Posterson, Gatth
APPLICANT: Posterson, Gatth
APPLICANT: Morris, Jason
APPLICANT: Morris, Jason
APPLICANT: Tissenbaum, Heidi
APPLICANT: Morris, Jason
APPLICANT: Koweek, Allison
TITLE OF INVENTION: IMPAIRED GLUCOSE TOLERANCE CONDITIONS
TITLE OF INVENTION: IMPAIRED GLUCOSE TOLERANCE CONDITIONS
FILE REFERENCE: OO786/351001
CURRENT APPLICATION NUMBER: US/08/857,076C
CURRENT FILING DATE: 1997-05-15
NUMBER OF SEQ ID NOS: 114
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Patent No. 6861256
GENERAL INCORMATION:
APPLICANT: Ruckun, Gary
APPLICANT: OGG, SCOTE
TITLE OF INVENTION: IMPRARED GLUCOSE TOLERANCE CONDITIONS
TITLE OF INVENTION: IMPAIRED GLUCOSE TOLERANCE CONDITIONS
TITLE OF INVENTION: IMPAIRED GLUCOSE TOLERANCE CONDITIONS
TITLE OF INVENTION: IMPAIRED GLUCOSE TOLERANCE CONDITIONS
FILE REFERENCE: 00786/351004
CURRENT APPLICATION NUMBER: 08/057,076
EARLIER PILLING DATE: 1997-05-15
EARLIER PILLING DATE: 1997-07-07
EARLIER PILLING DATE: 1997-07-07
EARLIER PILLING DATE: 1998-05-15
NUMBER OF SEQ ID NOS: 328
SOFTWARE: FREUESE FILLING DATE: 1998-05-15
NUMBER OF SEQ ID NOS: 328
SEQ ID NO 75
LENGTH: 112
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100.0%; Pred. No. 45;
ive 0; Mismatches 0; Indels
                                                                                                                                                            Length 105;
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100.0%; Pred. No. 42;
iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 75, Application US/08857076C Patent No. 6225120
                                                                  ; TYPE: PRT
; ORGANISM: Streptococcus pneumoniae
US-09-583-110-5144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Caenorhabditis elegans
                                                                                                                                                                            Best Local Similarity 100.
Matches 6; Conservative
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NUMBER OF SEQ ID NOS:
SEQ ID NO 5144
LENGTH: 105
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Best Local Similarity
Matches 6; Conserv
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28 AELEKE 33
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US-08-857-076-75
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US-09-205-658-75
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LENGTH: 112
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Length 121;

DB 2;

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Length 135;
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100.0%; Pred. No. 54;
tive 0; Mismatches
                             COMPUTER: CL/ACM ISOSOGO
COMPUTER: CL/ACM ISOSOGO
COMPUTER: CL/ACM ISOSOGO
COMPARE: CONKNOWN>
SOFTWARE: CL/ACM CONTO
APPLICATION DATA:
APPLICATION NUMBER: US/09/107,433
FILLING DATE: 30-Jun-1998
PRIOR APPLICATION NUMBER: 60/085131
PILLING DATE: May 12, 1998
APPLICATION NUMBER: 60/051553
ATTORNEY/AGENT INFORMATION:
NAME: Ariniello, Pamela Deneke
REGISTRATION NUMBER: 40,489
REGISTRATION NUMBER: 40,489
REGISTRANCE/DOCKET NUMBER: GTC-011
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc feature
LOCATION: (B) LOCATION 1...135
SEQUENCE DESCRIPTION: SEQ ID NO: 5156:
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Job time : 25.5069 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 135 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPAX: (781)893-8277
INFORMATION FOR SEQ ID NO: 5156:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: YES
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6; Conservative
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Best Local Similarity
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APPLICANT: Gary Breton et. al

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.2004001
CURRENT APPLICATION NUMBER: US/09/489,039A
CURRENT PILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR PILING DATE: 1999-01-29
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US-09-107-433-5156
i. Sequence 5156, Application US/09107433
i. Sequence 5156, Application
i. General No. 6800744
i. GENERAL INFORMATION:
i. APPLICANT: Lynn A Doucette-Stamm and David Bush
i. APPLICANT: Lynn A Doucette-Stamm SEQUENCES RELATING TO STREPTOCOCCUS PNEUMONIAE
SEQUENCES RELATING TO STREPTOCOCCUS PNEUMONIAE
TITLE OF INVENTION: SEQUENCES RELATING TO STREPTOCOCCUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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CORRESPONDENCE ADDRESS:
ADDRESSEE: GENOME THERAPEUTICS CORPORATION
STREET: 100 Beaver Street
CITY: Waltham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20.7%; Score 6; DB 1;
100.0%; Pred. No. 49;
tive 0; Mismatches
                                                                                                                     07917/025001
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Patent No. 6610836
TTURNEY/AGENT INPORMATION:
NAME: Fasse, J. Peter
NAME: Fasse, J. Peter
REGISTRATION NUMBER: 32,983
REFERENCE/DOCKET NUMBER: 07917
TELECOMMUNICATION INFORMATION:
TELEFAX: 617-542-5070
TELEFAX: 617-542-5070
TELEFAX: 617-542-8906
INFORMATION FOR SEQ ID NO: 12: SEQUENCE CHARACTERISTICS:
LENGTH: 124 amino acids
TYPE: amin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TYPE: PRT
; ORGANISM: Klebsiella pneumoniae
US-09-489-039A-10568
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: Massachusetts
COUNTRY: USA
ZIP: 02354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 10568
LENGTH: 132
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Best Local Similarity 100.
Matches 6; Conservative
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Best Local Similarity 100.
Matches 6; Conservative
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| US-09-252-991A-2977;<br>US-09-712-363-253<br>US-09-252-991A-1849;<br>US-09-902-540-15276<br>US-09-252-991A-2739 | 3 5 5                                   | 5 5  | b                  | ă                                       | US-09-583-110-5168<br>US-09-248-796A-1419 | Ď.                | US-09-252-991A-2569<br>US-09-543-681A-4223 | 5                                      | Ď.                | őő              | 5                                     | ŏ                | Ď:               | ã Ě                 | 5 <u>∺</u>        | ő  | Ď  | Ď                   | Ď                    | 5 E                 | 5 5                 | US-09-410-372-8                           | Ď.                | US-09-949-016-1011  | US-07-603-133B-30     | ö                    | 5                  | US-09-358-383C-22   | US-09-232-991A-294   | US-09-949-016-11458 | ä                   | <b>:</b>             | 5 E                 | Ď  | S                   | Ď ‡                            | 5 🛱               | Þ   | Þ                    | <b>;</b>           | Þ ;                  | בָּ כֹ               | US-09-252-991A-2705 | Þ                   | US-09-252-991A-3196 | Þ                    | Þ:              | <b>-</b> :       | US-09-540-U14-9     | US-US-252-991A-2109  | US-03-252-331A-253   | US-09-252-991A-2538 | US-09-270-767-42655                | US-10-091-841A-9    |
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| 17.2<br>17.2<br>17.2<br>17.2<br>3   | 100                                     | 120  | 2.5                | 2.5                                     | 2.5                                       | 7.2               | 2.5  | 7.2                                    | 7.                | 7. 7.           | ! ??                                  | .2               | 7.               | vi c                | , c               | ! ?!   | . ~!                                       | .2                  | 7.                   | vi c                |                     | . ~                                       | 2.0               | uj c                |                       | ! ??                 | 7                  | ٠.<br>د             | i c                  | . r.                | 7.                  | ~ .                  |                     | 7.2  | 7.2                 | vi c                           | , r               | 2.7   | 7.2                  | 7.2                | 7.2                  | . r                  | 4.0                 | 7.2                 | 7.2                 | 7.2                  | 7.2             | 2.2              | 2.5                 | vi c                 | i.c                  | , c                 | • ^                                |                     |
| ហលលលល   | ւտւ                                     | , w r  | ı CO L             | n w                                     | ഗഗ  | ı,                | n n  | ט נ                                    | Ŋ.                | ഹ ഹ             | 'n                                    | S                | ι Oτ             | ın u                | n u               | 'n   | o.   | S                   | S)                   | ın ı                | n u                 | n<br>n                                    | ιΩ                | n u                 | n n                   | υ<br>Ω               | 5                  | ı,                  | Λu                   | n LC                | S                   | ហ                    | տ տ                 | n m  | 'n                  | ı<br>G                         | ח ת               | 'n  | ហ                    | S                  | ហ                    | ın ı                 | ח ער                | 'n                  | S                   | 2                    | ហ               | LO I             | ın ı                | חי                   | n u                  | n u                 | יין ה                              | , <b>L</b>          |
| 909<br>906<br>906<br>708<br>808   | ) 00 0<br>00 0                          | 911  | 913                | 915                                     | 916                                       | 918               | 919  | 921                                    | 922               | 923             | 925                                   | 926              | 927              | 928                 | 929               | 931  | 932  | 933                 | 934                  | 935                 | 936                 | 938                                       | 939               | 940                 | 947                   | 943                  | 944                | 945                 | 946                  | 948                 | 949                 | 950                  | 951                 | 953  | 954                 | 955                            | 900               | 928   | 959                  | 096                | 1961                 | 962                  | 200                 | 196                 | 996                 | 196                  | 896             | 696              | 970                 | 971                  | 9/2<br>6/2           | 67.8                | 975                                | 926                 |
| Sequence 16397, A<br>Sequence 8, Appli<br>Sequence 172, App<br>Sequence 8, Appli<br>Sequence 8, Appli           | Sequence Jir, Appl<br>Sequence 10, Appl | Sequence 10314, A<br>Sequence 10314, A       | Sequence 158, App  | Sequence 10, Appr<br>Sequence 14104, A  | Sequence 6, Appli                         | Sequence 568, App | Sequence 11003, A                          | sequence sos, App<br>Sequence 8, Appli | Sequence 8, Appli | α, α            | . 51                                  | 15,              | 15,              | 1293                | 15,               | 4454   | 34274                                      | 49491               | 23835                | 33657               | 48874<br>9345       | 6797                                      | Sequence 49, Appl | 12624               | 12, <i>f</i><br>23764 | 25948,               | 6760,              | 44951,              | 32745,               | 11 4                | 10245,              | 29453,               | 7935,               | 31978.                                     | 11527,              | 2, Apr                         | Sequence 4, Appli | Sequence 11112. A                             | Sequence 19241, A    | 9958, A            | 23091,               | 2996                 | 117                 | 176                 | . 4                 | Sequence 16918, A    | 4, 1            | 4                | H.                  | ä                    | 2                    | · 1                 | 'nς                                |                     |
| US-09-902-540-16397<br>US-09-218-363-8<br>US-09-716-964B-172<br>US-09-772-105-8                                 | US-10-104-04/-5114<br>US-09-218-363-10  | US-09-232-391A-22631<br>US-09-489-039A-10314 | US-09-602-777A-158 | US-09-//2-105-10<br>US-09-902-540-14104 | US-09-408-020-6                           | US-09-205-258-568 | US-09-949-016-11003                        | US-10-004-860-568<br>US-08-393-889-8   | US-09-119-900-8   | US-08-472-630-8 | FCI-0594-09450-8<br>IIS-08-393-889-15 | US-09-119-900-15 | US-08-472-630-15 | US-09-902-540-12933 | PCT-US94-09450-15 | US-UY-Z5Z-YYLA-Z1634<br>TIS-00-134-0000-4454 | US-09-134-000C-4454<br>US-09-270-767-34274 | US-09-270-767-49491 | US-09-252-991A-23839 | US-09-270-767-33657 | US-09-270-767-48874 | US-U9-469-U39A-9343<br>HS-09-328-352-6797 | US-10-159-901-49  | US-09-902-540-12624 | US-09-890-813-12      | US-09-232-991A-25948 | US-09-328-352-6760 | US-09-270-767-44951 | US-09-252-991A-32745 | US-09-134-000C-5059 | US-09-949-016-10245 | US-09-252-991A-29453 | US-09-543-681A-7935 | US-US-343-661A-6131<br>TG-09-270-767-31978 | US-09-902-540-11527 | US-09-078-091-2                | US-09-078-091-4   | US-US-232-331A-23834<br>IIS-09-489-039A-11112 | US-09-248-796A-19241 | US-09-902-540-9958 | US-09-252-991A-23091 | US-09-252-991A-29966 | US-09-902-540-11785 | US-US-302-340-13231 | US-09-361-0368-1    | US-09-248-796A-16918 | US-09-772-105-4 | US-08-531-927B-4 | US-09-557-262-15    | US-09-489-039A-12694 | US-09-252-991A-22068 | US-08-411-607A-2    | US-U9-361-741-2<br>US-09-461-419-2 | US-09-949-016-10678 |
| 00000   | 900                                     | 4 (4 (                                       | 4 (7)              | <i>N</i> (4)                            | 9 0                                       | 1 (1              | 0 0  | N (7                                   | 10                | 0 5             | 4 (                                   | 1 (1             | 0                | 0                   | 4, (              | 40   | 4 (  | 1 (7)               | 0                    | N                   | 0 0                 | 4 0                                       | ~                 | 0                   | 0 0                   | A (                  | 1 (1               | 7                   | 0                    | 0 C                 | 1 (1                | 7                    | 0                   | 4 C  | 1 (4                | N                              | (1)               | 4 0   | 10                   | ~                  | ~                    | 7                    | N C                 | <b>v</b> (          | 4 C                 | 1 (1                 | N               | Н                | N                   | CI I                 | α,                   | (                   |                                    | N (2)               |
| 00000   | ب بو ہ                                  | 566  | 78                 | 9 6                                     | 56  | 272               |  |  |                   |                 |                                       |                  |                  |                     |                   |  |  |                     |                      |                     |                     |   |                   |                     |                       |                      |                    |                     |                      |                     |                     |                      |                     |  |                     |                                |                   |   |                      |                    |                      |                      |                     |                     |                     |                      | 25              | 25               | 25                  | 55                   | 5 6                  |                     |                                    |                     |
| 7 5 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6   |   |  |                    |   |   |                   |  |  |                   |                 |                                       |                  |                  |                     |                   |  |  | ·N                  | (V                   | IN                  | · v · c             | • 0                                       |                   |                     | ' A C                 | 10                   |                    | ď                   | .4 (                 | u, c                |                     | "                    |                     |  | 4                   |                                |                   |   | ,                    |                    |                      |                      |                     |                     |                     | 4 ^4                 | ~               | ď                | a                   | o.                   | o .                  | O 4                 |                                    |                     |
| 17.2 266<br>17.2 266<br>17.2 266<br>17.2 266  | N 01 C                                  | 17.5   | 17.2               | 17.2                                    | 17.2                                      | 17.2              | 17.2                                       | 17.2                                   | 17.2              | 17.2            | 17.7                                  | 17.2             | 17.2             | 17.2                | 17.2              | 17.  | 12   | 17.                 | 17.                  | 17.                 | 17.                 | 12  | 17.               | 17.                 | 17.                   | 1 -                  | 11                 | 17                  | 17                   | 1 -                 | 17.                 | 17                   | 17.                 | 11   | 17.                 | 17.                            | 17.               | 1.  |                      | 17.                | 17.                  | 17.                  | 17.                 | 7                   | 1.                  | 17.2                 | 17.2            | 17.2             | 17.2                | 17.2                 | 17.2                 | 17.2                | 17.                                | 11                  |
| 00000   | 17.2                                    | 11:  | 17.                | 17.                                     | 17.                                       | 17.               | 17.  | 17.                                    | 17.               | 17.             | 17.                                   | 17.              | 17.              | 17.                 | 17.               | 17.  | 17.  | 17.                 | 17.                  | 17.                 | 17.                 | 1.  | 17.               | 17.                 | 17.                   |                      |                    |                     | 17                   | 17                  | 17                  | 17                   | 17                  | 11   | 17.                 | 17.                            | 17                | 1.  | 1.                   | 17.                | 17.                  | 17                   | 17                  | 7,                  | 1.                  | 17                   | 17              | 17               | 17                  | 17                   | 17                   | 17                  | 17                                 |                     |

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100.08;
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Best Local Similarity 100.
Matches 9; Conservative
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RAAFLEK 76
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                                                                                                                                TYPE: PRT
ORGANISM: Human
                                                                                                                                                       US-09-949-016-11210
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Patent No. 681239

GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CLO01307

CURRENT APPLICATION NUMBER: 00/29149,016

CURRENT APPLICATION NUMBER: 60/241,755

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR APPLICATION NUMBER: 60/237,768

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-09-08

NUMBER: 60/231,498

NUMBER OF SEQ ID NOS: 207012

SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 2
US-09-949-016-11210
i Sequence 11210, Application US/09949016
j Sequence 11210, Application US/09949016
j Edtent No. 6812339
j GENERAL INFORMATION:
j APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
j FILE REFERENCE: CL001107
j CURRENT APPLICATION NUMBER: US/09/949,016
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                                                                                                                     582, App
2, Appli
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28498, A
12986, A
8, Appli
8, Appli
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32, Appl
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Sequence 8
Sequence 8
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                       ALIGNMENTS
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100.0%; Pre-
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Best Local Similarity 100.
Matches 14; Conservative
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ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                     US-09-949-016-10895
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Sequence 4130, Application US/09107532A
| Patent No. 6583275
| GENERAL INFORMATION:
| TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
| RYPEROCOCCUS FAECTUM FOR DIAGNOSTICS AND THERAPEUTICS
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                                  Length 35;
                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 7310
CORRESPONDENCE BADRESS:
ADDRESSEE: GENOME THERAPEUTICS CORPORATION
STREET: 100 Beaver Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT TYPE: C., COMPUTER: PC
COMPUTER: PC
COMPUTER: PC
COMPUTER: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/107,532A
FILING DATE: 30-Jun-1998
PRIOR APPLICATION NUMBER: 60/085,598
FILING DATE: 14 May 1998
APPLICATION NUMBER: 60/051571
FILING DATE: JUNY 2, 1997
ATORNEY/AGENT INFORMATION:
NAME: Ariniello, Pamela Deneke
REGISTRATION NUMBER: 40,489
REBERBURGE/BOCKET NUMBER: GTC-012
TELECOMMUNICATION INFORMATION:
TELEFHONE: (781)893-5007
TELEFHONE: (781)893-8277
INFORMATION FOR SEQ ID NO: 4130:
SEQUENCE CHARACTERISTICS:
LIENGTH: 86 amino acid
TYPE: amino acid
TYPE: amino acid
TYPE: amino acid
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100.0%; Pred. No. 57;
tive 0; Mismatches
                              Query Match 20.7%; Score 6; DB 2; Best Local Similarity 100.0%; Pred. No. 26; Matches 6; Conservative 0; Mismatches
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; LOCATION: (B) LOCATION 1...86
; SEQUENCE DESCRIPTION: SEQ ID NO: 4130:
US-09-107-532A-4130
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 02354
COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660
                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Waltham
STATE: Massachusetts
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: YES
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Best Local Similarity 100.
Matches 6; Conservative
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US-09-107-532A-4130
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US-09-902-540-13617
US-10-029-217A-20
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APPLICANT: Milan, Aileen
APPLICANT: Burk, Mark J.
APPLICANT: Short, Jay M.
TITLE OF INVENTION: EXPOXIDE HYDROLASES, NUCLEIC ACIDS ENCODING THEM AND METHODS
TITLE OF INVENTION: OF MAKING AND USING THEM
FILE REFERENCE: 09010-031001
CURRENT APPLICATION NUMBER: US 10/214,473
PRIOR APPLICATION NUMBER: US 60/309,478
PRIOR APPLICATION NUMBER: US 60/309,478
PRIOR PILING DATE: 2001-08-03
PRIOR FILING DATE: 2001-08-03
PRIOR FILING DATE: 2001-09-03
PRIOR FILING DATE: 2001-09-03
SEQ ID NO 66
SEQ ID NO 66
LENGTH: 401
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                 Length 375;
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24.1%; Score 7; DB 2;
Best Local Similarity 100.0%; Pred. No. 22;
Matches 7; Conservative 0; Mismatches
                 DB 2;
               Query Match 24.1%; Score 7; DB 2
Best Local Similarity 100.0%; Pred. No. 21;
Matches 7; Conservative 0; Mismatches
                                                                                                                                                                                                                     ; Sequence 66, Application US/10272490 ; Patent No. 6943001
                                                                                                                                                                                                                                                                         APPLICANT: Zhao, Lishan
APPLICANT: Mathur, Eric J.
APPLICANT: Weiner, David
APPLICANT: Richardson, Toby
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28 AELRKRV 34
                                                                                                                          2 RTRVAEL 8
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ORGANISM: Unknown
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US-10-029-217A-20
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APPLICANT: Gary Breton et. al
APPLICANT: Gary Breton et. al
APPLICANT: Gary Breton et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: PREUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.2004001
CURRENT FELLNG DATE: 2000-01-27
CURRENT FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 11896
LENGTH: 141
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
  ADDRESSEE: GENOME THERAPEUTICS CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 2;
                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/107,433
PILING DATE: 30-Jun-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/085131
PILING DATE: May 12, 1998
APPLICATION NUMBER: 60/05153
FILING DATE: July 2, 1997
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Streptococcus pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
20.7%; Score 6; DB 2
Best Local Similarity 100.0%; Pred. No. 90;
Matches 6; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature; LOCATION: (B) LOCATION 1...135; SEQUENCE DESCRIPTION: SEQ ID NO: 5156: US-09-107-433-5156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: GTC-011
                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Ariniello, Pamela Deneke
REGISTRATION NUMBER: 40,489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20.7%; Score 6; I
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 11896, Application US/09489039A; Patent No. 6610836
                                                                                                                                                    MEDIUM TYPE: CD/ROM ISO9660
COMPUTER: «Unknown»
OPERATING SYSTEM: «Unknown»
SOFTWARE: «Unknown»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: (781)893-5007
                    STREET: 100 Beaver Street
CITY: Waltham
STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 135 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 5156:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (781)893-8277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Klebsiella pneumoniae
                                                                                                           ZIP: 02354
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: YES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.
Matches 6; Conservative
                                                                                           COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8 ENTALR 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-489-039A-11896
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APPLICANT: Lynn Doucette-Stamm et al.

APPLICANT: Lynn Doucette-Stamm et al.

TITLE OF INVENTION: Nacleic Acid and Amino Acid Sequences Relating to Streptococcus
TITLE OF INVENTION: Nacleic Acid and Amino Acid Sequences Relating to Streptococcus
TITLE OF INVENTION: Nacleic Acid and Therapeutics

FILE REFERENCE: PATH00-07A
CURRENT APPLICATION NUMBER: US/09/583,110

CURRENT APPLICATION NUMBER: US 60/085,131

PRIOR PILING DATE: 1998-06-12

PRIOR APPLICATION NUMBER: US 60/051,553

PRIOR APPLICATION NUMBER: US 60/051,553

PRIOR PILING DATE: 1997-07-02
NUMBER OF SEQ ID NOS: 5322

SEQ ID NO 5144

LENGTH: 105
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID
SEQUENCES RELATING TO STREPTOCOCCUS PNEUMONIAE
THERAPEUTICS
NUMBER OF SEQUENCES: 5206
CORRESPONDENCE ADDRESS:
                                                           APPLICANT: Goldman, Barry S.
APPLICANT: Goldman, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Steven C.
APPLICANT: Steven C.
APPLICANT: Wiegand, Roger C.
TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof FILE REFERENCE: 38-10(15849)B
CURRENT APPLICATION NUMBER: US/09/902,540
CURRENT FILING DATE: 2001-07-10
PRIOR APPLICATION NUMBER: 60/217,883
PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 13617
LENGTH: 92
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 92;
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100.0%; Pred. No. 69;
iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 6; DB 2;
; Pred. No. 61;
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100.0%; Pred. No. c...
0; Mismatches
Sequence 13617, Application US/09902540 Patent No. 6833447 GENERAL INFORMATION:
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Patent No. 6699703
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US-09-107-433-5156
; Sequence 5156, Application US/09107433
; Patent No. 6800744
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TYPE: PRT
; ORGANISM: Streptococcus pneumoniae
US-09-583-110-5144
                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT ORGANISM: Myxococcus xanthus
                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 100.
Matches 6; Conservative
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Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                   US-09-902-540-13617
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Sequence 626, Application US/09732210

Sequence 626, Application US/09732210

Patent No. 6573361

GENERAL INPORMATION:

APPLICANT: Bunkers, Greg J.

APPLICANT: Liang, Jihong

APPLICANT: Mittanck, Gindy A.

APPLICANT: Mittanck, Gindy A.

APPLICANT: Man, Younie S.

TITLE OF INVENTION: Anti-fungal Proteins and Methods for Their Use

FILE REFERENCE: 38-21(15036)B

CURRENT APPLICATION NUMBER: US/09/732,210

CURRENT PILING DATE: 1999-12-07

PRIOR APPLICATION NUMBER: US 60/169,513

PRIOR APPLICATION NUMBER: US 60/169,340

PRIOR PRILING DATE: 1999-12-07

NUMBER OF SEQ ID NOS: 1753

SEQ ID NO 626

SEQ ID NO 626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Pizza, Mariagrazia
APPLICANT: Grandi, Guido
TITLE OF INVENTION: Neisserial Antigens
FILE REPERENCE: CHIROLGO
CURRENT APPLICATION NUMBER: US/09/303,518D
CURRENT FILING DATE: 1999-04-30
NUMBER OF SEQ ID NOS: 1098
SOFTWARE: Patentin version 3.1
SEQ ID NO 826
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100.0%; Pred. No. 95;
stive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Neisseria gonorrhoeae
US-09-303-518D-826
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 100.
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Best Local Similarity 100.
Matches 6; Conservative
Rappuoli, Rino
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16 RKRVGR 21
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                                                                                                                                                                                                                              Sequence 3869, Application US/09583110

Fatent No. 6699703

GENERAL INFORMATION:
FATEL OF INVENTION:
TITLE OF INVENTION:
FILE REFERENCE:
FILE OF INVENTION:
FILE REFERENCE:

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Patent No. 6562958
GENERAL INPORMATION:
GENERAL INPORMATION:
GENERAL INPORMATION:
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GT-99-03PA
CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT PILING DATE: 1999-06-04
NUMBER OF SEQ ID NOS: 8252
SEQ ID NO 6345
LENGTH: 142
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APPLICANT: Scarlato, Vincenzo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT CORGANISM: Streptococcus pneumoniae US-09-583-110-3869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TYPE: PRT
; ORGANISM: Acinetobacter baumannii
US-09-328-352-6345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Masignani, Vega
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 20.7
Best Local Similarity 100.
Matches 6; Conservative
                                   12 LRTRVA 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4 FLEKEN 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 AFLEKE 8
                                                                                             4 LRTRVA 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 13
US-09-328-352-6345
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Sequence 303, App
Sequence 303, App
Sequence 24950, A
Sequence 8632, Ap
Sequence 1965, A
Sequence 13021, A
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7580, Ap
13, Appl
14685, A
16333, A
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13239, A
9985, Ap
31704, A
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12636, A
6781, Ap
320, App
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24, Appl
7090, Ap
14097, A
18540, A
                                         2, Appli
7298, Ap
5, Appli
5, Appli
1, Appli
6, Appli
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10575, A
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| Sequence 2, Appli<br>Sequence 2, Appli   | 00000   | equence 2,<br>equence 2,<br>equence 2,<br>equence 2,  | equence 2,<br>equence 2,<br>equence 28,  | equence 28,<br>equence 2,<br>equence 3,  | equence 28,<br>equence 133<br>equence 577  | equence 32  | equence 19,<br>equence 2,<br>equence 109  | equence 238<br>equence 5,  | 9 0 0   | equence 328<br>equence 707<br>equence 169  | equence 428<br>equence 682  | e 411,<br>e 932,<br>e 1486  | equence 11<br>equence 97   | equence 48,<br>equence 48,  | equence 8067,  | equence 48,<br>equence 361,  | equence 363,<br>equence 367,                                 | equence 10<br>equence 10<br>equence 65  | equence 101,   | equence 46,                                 | equence 107,  | equence 8067                          | equence 129<br>equence 129  | equence 10447<br>equence 3909,   | equence 54,<br>equence 54,  | Sequence 54, Appl<br>Sequence 54, Appl<br>Sequence 54, Appl   |
|--|---|---|--|--|--|---|---|--|---|--|---|---|--|---|--|--|--|---|--|---|---|---------------------------------------|---|--|---|---|
| 1 US-08-508-735-2<br>1 US-08-627-610-2<br>1 US-08-306-511A-2<br>1 US-08-893-274-2<br>1 US-08-848-251-2<br>1 US-08-486-047-2<br>1 US-08-120-130-2   | s s s s   | 8888  | ds<br>ds   | si si  | Sin Sin  | Sin   | P.S.  | -SU<br>FO  | s n   | sn ns  | us-   | s s   | Sn s   | sn<br>ns  | sn<br>ns   | Sn Sn  | sn s   | SD S  | ns-  | US  | SD  | s s                                   | s s   | s s  | -sn   | us<br>us  |
| 1126<br>1156<br>1156   | 156<br>156<br>156<br>156  | 156<br>156<br>156<br>156  | 156<br>156<br>156  | 156<br>156<br>156  | 156<br>156   | 156   | 156   | 157  | 158<br>160  | 160<br>160<br>161  | 161   | 161<br>161  | 162  | 166<br>166  | 166<br>166   | 166  | 167  | 167   | 167  | 167   | 167   | 168                                   | 168   | 169  | 170   | 170<br>170<br>170   |
| 111111111111111111111111111111111111111  | 11.6<br>11.6<br>11.6<br>11.6  | 11.6<br>11.6<br>11.6  | 11.6<br>11.6<br>11.6   | 11.6<br>11.6<br>11.6   | 11.6   | 11.6  | 11.6  | 11.6   | 11.6<br>11.6  | 11.6<br>11.6<br>11.6   | 11.6  | 11.6<br>11.6<br>11.6  | 11.6   | 11.6  | 11.6   | 11.6   | 11.6   | 11.6  | 11.6   | 11.6  | 11.6  | 11.6                                  | 11.6  | 11.6<br>11.6   | 11.6<br>11.6  | 11.6<br>11.6<br>11.6  |
| លលល់លំបំបំព  | ហលលលល   | ហហហហ  | ហហហ  | വവവ  | សសល  | ហេហ   | ո տ տ   | ហេស  | ոտտ   | ហលល  | ហហ  | ո տ տ   | ហហហ  | ດເນ   | ហហ   | ហហ   | ហហរ  | ռտտ   | സസ   | rU n  | ոտտ   | ומוח                                  | υn  | വവ   | ហហ  | ហហហ   |
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| Sequence 478, App<br>Sequence 287, App<br>Sequence 5156, Ap<br>Sequence 6, Appli<br>Sequence 6, Appli<br>Sequence 4, Appli<br>Sequence 4, Appli  | Sequence 4,<br>Sequence 111<br>Sequence 16,<br>Sequence 16,<br>Sequence 16,   |   | Sequence 25,<br>Sequence 16,<br>Sequence 4,  | Sequence 16,<br>Sequence 16,<br>Sequence 16,   | Sequence 16,<br>Sequence 16,<br>Sequence 4.  | Sequence 16,  | Sequence 425<br>Sequence 286  | Sequence 4, Aps Sequence 42175   | Sequence 46853<br>Sequence 33, P  | Sequence 114,<br>Sequence 16687<br>Sequence 11283  | Sequence 366,<br>Sequence 369,  |   | Sequence 109,<br>Sequence 10021  | Sequence  | Sequence 25434<br>Sequence 26454   | Sequence   | Sequenci   | Sequence  | Sequence 987<br>Sequence 826                                     | Sequence 24                                 | Sequence 654,<br>Sequence 14, A   | Sequence 2,                           | Sequence 2,<br>Sequence 2,  | Sequence 42<br>Sequence 16   | Sequence 682<br>Sequence 642  | Sequence 2, Appl<br>Sequence 2, Appl<br>Sequence 2, Appl  |
| Sequence 478, Ap<br>Sequence 287, Ap<br>Sequence 5156, A<br>Sequence 41376, A<br>Sequence 6, Appl<br>Sequence 4, Appl<br>Sequence 4, Appl<br>Sequence 4, Appl  | Sequence 4,<br>Sequence 111<br>Sequence 16,<br>Sequence 16,<br>Sequence 16,   | Sequence 16,<br>Sequence 4,<br>Sequence 16,<br>Sequence 16,   | Sequence 25,<br>Sequence 16,<br>Sequence 4,  | Sequence 16,<br>Sequence 16,<br>Sequence 16,   | Sequence 16,<br>Sequence 16,<br>Sequence 4.  | Sequence 16,  | Sequence 4, Ab<br>Sequence 4254,<br>Sequence 286,   | Sequence 4, Aps Sequence 42175   | Sequence 46853<br>Sequence 33, P  | Sequence 114,<br>Sequence 16687<br>Sequence 11283  | Sequence 366,<br>Sequence 369,  | Sequence 109,<br>Sequence 109,<br>Sequence 106,   | Sequence 109,<br>Sequence 10021  | Sequence  | Sequence   | Sequence   | Sequenci   | Sequence  | Sequence 98<br>Sequence 82                                       | Sequence 24                                 | Sequence 65<br>Sequence 14  | Sequence 2,                           | Sequence 2,<br>Sequence 2,  | Sequence 42<br>Sequence 16   | Sequence 6826, A<br>Sequence 642, Ap  | Sequence 2, Appl<br>Sequence 2, Appl<br>Sequence 2, Appl  |
| 2 US-09-830-230A-478 Sequence 478, Appl 2 US-09-615-192A-287 Sequence 287, Appl 2 US-09-612-976-4376 Sequence 5156, Appl 2 US-08-306-512-6 Sequence 6, Appl 1 US-08-306-511A-4 Sequence 4, Appl 1 US-08-3274-4 Sequence 4, Appl 2 US-08-3507-4 Sequence 4, App | 2 US-09-016-750C-4 Sequence 4, 28-09-016-750C-4 Sequence 111 US-08-474-177-16 Sequence 15, 1 US-08-487-033-16 Sequence 16, 1 US-08-480-810-16 Sequence 16,  | 1 US-08-508-735-16 Sequence 16, 1 US-08-627-610-4 Sequence 4, 1 US-08-848-251-16 Sequence 16, 1 US-08-486-047-16 Sequence 16,                                     | 2 US-08-384-106A-25 Sequence 25,<br>2 US-09-120-130-16 Sequence 16,<br>2 US-08-581-918A-4 Sequence 4,                            | 2 US-09-115-252-16 Sequence 16,<br>2 US-08-986-515-16 Sequence 16,<br>2 US-09-120-128-16 Sequence 16,                      | 2 US-09-120-129-16 Sequence 16, 2 US-09-201-139-16 Sequence 16, 2 US-08-346-147B-4 Sequence 4.                             | 2 US-09-120-131-16 Sequence 16, 2 US-08-2-936-4 Sequence 4, 2 US-08-407-2140-4 Sequence 4, 2 US-08-407-2140-4   | 2 US-09-134-001C-4254 Sequence 286,4,2 US-09-605-703B-286 Sequence 286,4                  | 4 PCT-US95-04636-4 Sequence 4, Ag<br>2 US-09-270-757-42179 Sequence 42172<br>2 US-00-270-540-11415 Sequence 42172                            | 2 US-09-270-767-46853 Sequence 46852 Sequence 33, P. Sequence 34, P. Sequence 35, P. Sequence | 2 US-09-345-236B-114 Sequence 114,<br>2 US-09-252-91A-1687 Sequence 16687<br>2 US-09-902-540-11283 Sequence 11283                              | 2 US-08-513-974B-366 Sequence 366, 2 US-08-513-974B-369 Sequence 369,   | 2 US-08-776-971-109 Sequence 109, 2 US-08-776-971-109 Sequence 109, 2 US-09-576-290-106 | 2 US-09-576-280-109 Sequence 109, 2 US-09-902-840-10021 Sequence 10021 2 US-09-716-147-106 Sequence 106                                    | 2 US-09-716-147-109 Sequence<br>2 US-09-266-965-121 Sequence                | 2 US-09-252-991A-25434 Sequence<br>2 US-09-252-991A-26454 Sequence                   | 2 US-09-252-991A-27269 Sequence<br>1 US-08-154-915-4 Sequence                | 2 US-08-384-106A-16 Sequence<br>2 US-08-384-106A-24 Sequence | 2 US-09-0/5-505-2<br>2 US-09-107-532A-6636 Sequenc<br>4 PCT-11893-09945-4 Semienc               | 2 US-10-101-464A-987 Sequence 982 US-09-303-518D-826 Sequence 82 | 2 US-09-252-991A-24718 Sequence 24          | 2 US-10-464-939-14 Sequence 19 2 US-10-464-939-14 Sequence 14 2 US-10-466-7038-1748 Seminance 17                                    | 1 US-08-446-908-2 Sequence 2,         | 1 US-08-231-205A-2 Sequence 2,<br>1 US-08-871-161-2 Sequence 2,                   | 2 US-09-621-976-4212 Sequence 42 US-09-248-796A-16580 Sequence 16                        | 2 US-09-513-999C-6826 Sequence 6826, A<br>2 US-09-640-211A-642 Sequence 642, Ap                   | 1 US-08-474-177-2 Sequence 2, Appl<br>1 US-08-487-033-2 Sequence 2, Appl<br>1 US-08-480-810-2 Sequence 2, Appl                            |
| US-09-830-230A-478 Sequence 478, Appl US-09-615-192A-287 Sequence 287, Appl US-09-107-433-5156 Sequence 5156, Appl US-09-621-976-4376 Sequence 6, Appl US-08-893-274-4 Sequence 4, Appl US-08-893-893-893-893-893-893-893-893-893-89  | 11.6 137 2 US-09-016-750C-4 Sequence 4, 11.6 138 1 US-08-474-177-16 Sequence 11.1 138 1 US-08-474-177-16 Sequence 16, 11.6 138 1 US-08-487-033-16 Sequence 16, 11.6 138 1 US-08-480-810-16 Sequence 16, 11.6 138 1 US-08-480-810-16 | 11.6 138 1 US-08-508-735-16 Sequence 16, 11.6 138 1 US-08-627-610-4 Sequence 4, 11.6 138 1 US-08-848-251-16 Sequence 16, 11.6 138 1 US-08-486-047-16 Sequence 16, | 11.6 138 2 US-08-384-106A-25 Sequence 25,<br>11.6 138 2 US-09-120-130-16 Sequence 16,<br>11.6 138 2 US-08-581-918A-4 Sequence 4, | 11.6 138 2 US-09-115-252-16 Sequence 16, 11.6 138 2 US-09-986-515-16 Sequence 16, 11.6 138 2 US-09-120-128-16 Sequence 16, | 11.6 138 2 US-09-120-129-16 Sequence 16, 11.6 138 2 US-09-2011-139-16 Sequence 16, 11.6 138 2 US-08-246-1478-4 Sequence 4. | 11.6 138 2 US-09-120-131-16 Sequence 16, 11.6 138 2 US-08-082-936-4 Sequence 4, 11.6 138 2 US-08-082-936-4 Sequence 4, 11.6 138 2 US-08-082-936-4 Sequence 4, 11.6 138 2 US-08-082-936-14 | 11.6 138 2 US-09-134-001C-4254 Sequence 4254, 11.6 138 2 US-09-605-703B-286 Sequence 286, | 11.6 138 4 PCT-US95-04636-4 Sequence 4, Ag<br>11.6 140 2 US-09-270-767-42179 Sequence 42172<br>11.6 141 2 US-09-200-200-11415 Sequence 42172 | 11.6 142 2 US-09-719-554-33 Sequence 33, P  | 11.6 143 2 US-09-345-236B-114 Sequence 114,<br>11.6 143 2 US-09-525-991A-16687 Sequence 16687<br>11.6 143 2 US-09-902-540-11283 Sequence 11283 | 11.6 144 2 US-08-513-974B-366 Sequence 366, 11.6 144 2 US-08-513-974B-369 Sequence 369, 11.6 144 2 US-08-513-974B-369 | 11.6 144 2 US-09-776-291-109 Sequence 109, 11.6 144 2 US-09-576-290-106 Sequence 106,   | 11.6 144 2 US-09-576-290-109 Sequence 109,<br>11.6 144 2 US-09-902-540-10021 Sequence 10021<br>11.6 144 2 US-09-902-540-106 Sequence 10021 | 11.6 144 2 US-09-716-144-109 Sequence 11.6 145 2 US-09-266-965-121 Sequence | 11.6 145 2 US-09-252-991A-25434 Sequenci<br>11.6 146 2 US-09-252-991A-26454 Sequenci | 11.6 146 2 US-09-252-991A-27269 Sequence 11.6 148 1 US-08-154-915-4 Sequence | 11.6 148 2 US-08-384-106A-16 Sequence                        | 11.6 148 2 US-U9-U7-505-2<br>11.6 148 2 US-U9-107-532A-636 Sequenc<br>11.6 148 4 PCT-US9-0945-4 | 11.6 149 2 US-10-101-464A-987 Sequence 98                        | 11.6 150 2 US-09-252-991A-24718 Sequence 24 | 11.6 151 2 US-103-04U-411A-054 Sequence 05<br>11.6 151 2 US-10-4464-939-14 Sequence 14<br>11.6 151 2 US-104-614-939-14 Seminance 14 | 11.6 154 1 US-08-46-908-2 Sequence 2, | 11.6 154 1 US-08-231-205A-2 Sequence 2,<br>11.6 154 1 US-08-871-161-2 Sequence 2, | 11.6 154 2 US-09-621-976-4212 Sequence 42<br>11.6 154 2 US-09-248-796A-16580 Sequence 16 | 11.6 154 2 US-09-513-999C-6826 Sequence 6826, A<br>11.6 155 2 US-09-640-211A-642 Sequence 642, Ap | 11.6 156 1 US-08-474-177-2 Sequence 2, Appl<br>11.6 156 1 US-08-487-033-2 Sequence 2, Appl<br>11.6 156 1 US-08-480-810-2 Sequence 2, Appl |

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| US-09-270-767-40384 US-09-270-767-56600 US-09-248-796A-24434 US-08-849-376-2 US-08-849-376-2 US-08-849-376-2 US-08-849-376-4 US-08-849-376-4 US-08-849-376-4 US-09-248-796A-15697 US-09-949-016-949 US-09-949-016-949 US-09-328-352-4681 US-09-328-352-4681 US-09-328-352-4681 US-09-328-352-4681 US-09-328-352-4681 US-09-328-352-4681 US-09-328-352-4681 US-09-328-352-4681 US-09-328-352-4681 US-09-18-475-523 US-09-18-778-791-22 US-09-18-778-791-22 US-09-18-778-791-22 US-09-18-778-791-22 US-09-252-991A-1144 US-09-248-796A-19139 US-09-248-796A-18199 US-09-248-796A-18199 US-09-252-991A-28198 US-09-248-796A-18199 US-09-252-991A-28199  |  |
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| Sequence 54, Appl Sequence 54, Appl Sequence 1243, A Sequence 51, Appl Sequence 61, Appl Sequence 61, Appl Sequence 62, Appl Sequence 62, Appl Sequence 62, Appl Sequence 61, Appl Sequence 62, Appl Sequence 62, Appl Sequence 63, Appl Sequence 11, Appl Sequence 296, Appl Sequence 206, Appl Sequence 296, Appl Sequence 206, Appl Sequence 207, Appl Sequence 2073, Appl Sequence  | , , , , , , , , , , , , , , , , , , ,  |
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| US-08-485-863A-54         Sequence           US-08-485-049D-54         Sequence           US-09-485-049A-12433         Sequence           US-09-772-719B-54         Sequence           US-09-770-767-45018         Sequence           US-09-270-767-3814         Sequence           US-09-107-5346-5272         Sequence           US-09-107-5340-1213         Sequence           US-09-107-5340-1223         Sequence           US-09-108-540-149-2         Sequence           US-08-609-9         Sequence           US-08-108-140-2         Sequence           US-08-141-139-2         Sequence           US-08-141-139-2         Sequence           US-08-141-139-2         Sequence           US-09-131-109-131         Sequence           US-09-131-109-131         Sequence           US-09-131-109-131         Sequence           US-09-131-109-131         Sequence           US-09-131-109-131   | 2 US-09-28-60-4-40<br>2 US-09-252-991A-26912 Sequence<br>2 US-09-270-767-31789 Sequence<br>2 US-09-270-767-60029 Sequence<br>2 US-09-252-991A-23063 Sequence<br>2 US-09-252-991A-23063 Sequence<br>2 US-09-489-033A-11511 Sequence<br>2 US-09-583-110-4224 Sequence<br>2 US-09-583-110-4224 Sequence<br>2 US-09-902-540-12260 Sequence<br>2 US-09-252-991A-25163 Sequence<br>2 US-09-902-540-1467 Sequence   |
| US-08-485-863A-54 US-08-485-049D-54 US-09-489-039A-12433 US-09-782-139-637A-359 US-09-199-637A-359 US-09-199-637A-359 US-09-199-637A-359 US-09-199-637A-359 US-09-199-637A-359 US-09-199-637A-359 US-09-199-637A-359 US-09-199-637A-359 US-09-199-637A-359 US-09-107-532A-5272 US-09-107-532A-5272 US-09-107-532A-5272 US-09-107-532A-591A-18339 US-08-252-991A-18339 US-08-252-991A-18339 US-08-261-1995-100-1995 US-09-252-991A-18339 US-09-252-991A-28915 US-09-252-991A-28918 US-09-252-991A-28918 US-09-252-991A-28938 US-09 | 11.6 194 2 US-08-284 Por A-491 Sequence 11.6 194 2 US-09-252-991A-26912 Sequence 11.6 194 2 US-09-270-767-31789 Sequence 11.6 194 2 US-09-270-767-313-2688 Sequence 11.6 196 2 US-09-107-4313-2688 Sequence 11.6 196 2 US-09-252-991A-23063 Sequence 11.6 196 2 US-09-252-991A-23063 Sequence 11.6 196 2 US-09-583-110-4224 Sequence 11.6 196 2 US-09-583-110-4224 Sequence 11.6 197 2 US-09-902-540-12260 Sequence 11.6 197 2 US-09-902-540-12260 Sequence 11.6 197 2 US-09-902-540-12260 Sequence 11.6 197 2 US-09-902-540-12467 Sequence 11.6 197 2 US-09-902-540-14467 Sequence  |

| Sequence 4, Appl:<br>Sequence 3, Appl:<br>Sequence 2, Appl:<br>Sequence 4, Appl:<br>Sequence 5181, Appl:<br>Sequence 2907. Appl:  | adnence   | equence<br>equence                       | equence  | equence  | equence   | equence                              | ednence                                | equence  | equence                                  | equence                                 | equence                                 | ednence                                | equence  | ednence                             | equence   | equence                                 | ednence                              | equence  | equence   | equence                               | ednence   | equence                              | edneuce                              | equence                                 | equence                                 | equence                                 | equence                                 | equence                                 | equence                                | equence 8586, A                     | equence 146                            | equence 876                             | equence 293<br>equence 950  | equence 43                              | equence                             | equence 56                           | equence 5,                          | equence 23                          | equence 58                          | equence 61'                          | equence 23                           |  |
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| US-09-776-976-4<br>US-09-506-852-3<br>US-09-909-547-2<br>US-09-909-547-4<br>US-09-134-0000-5181<br>US-09-540-336-2907   | US-10-392-706-3<br>US-09-724-864-51   |  | US-09-902-540-97<br>US-09-248-796A-1                                     | US-09-252-991A-2   | US-09-198-452A-1<br>US-09-457-568-22                                      |                                      |  |  | US-09-538-092-198                        |   |   | US-09-612-473-8                        | US-09-252-991A-1<br>US-09-252-991A-2                                     | US-09-489-039A-1010                 | US-09-270-767-61<br>US-09-919-039-93  |   | US-08-622-353-8                      |  |   |                                       |   |                                      |                                      |   |   |   |   |   |  |                                     |  |   | US-09-252-991A-2937<br>US-09-538-092-950                          |   |                                     |                                      |                                     |                                     |                                     |                                      |                                      |  |
| 247<br>247<br>247<br>247<br>247<br>247<br>247<br>247<br>247<br>247  |   |  |  |  |   |                                      |  |  |  |   |   |  |  |                                     |   |   |                                      |  |   |                                       |   |                                      |                                      |   |   |   |   |   |  |                                     |  |   |   |   |                                     |                                      |                                     |                                     |                                     |                                      |                                      |  |
| 55555   | 122   | 111                                      |  | 11   | 1 7 7   | 122                                  | 111                                    | i i  | 121                                      | 11.                                     | 11.                                     | 11                                     | 11.  | 11                                  | 11.   | 11:                                     | i i i                                | <u> </u>   | 11.   | 11.                                   | <u> </u>  | 11.                                  |                                      | 11.                                     | 11                                      | :<br>:<br>:                             | 11.                                     | 11:                                     | 1                                      | 1:                                  | 11                                     | 1:                                      | 1   | 11                                      | i:                                  | 1                                    | 11.                                 | 11.                                 | 11.                                 | 11                                   | 1:                                   | 11                                     |
| 758<br>759<br>761<br>762<br>763   | 764   | 767<br>768                               | 769<br>770   | 771  | 773   | 775                                  | 777                                    | 778  | 780                                      | 782                                     | 783                                     | 785                                    | 786  | 788                                 | 789<br>790  | 791                                     | 793                                  | 794<br>795   | 796   | 798                                   | 800   | 801                                  | 803                                  | 804                                     | 806                                     | 807                                     | 608                                     | 810                                     | 812                                    | 813                                 | 815                                    | 816                                     | 817   | 819                                     | 820                                 | 822                                  | 823                                 | 824                                 | 825<br>825                          | 827                                  | 828                                  | 830<br>830                             |
| O. et O. O. et  |   |  |  |  |   |                                      |  |  |  |   |   |  |  |                                     |   |   |                                      |  |   |                                       |   |                                      |                                      |   |   |   |   |   |  |                                     |  |   |   |   |                                     |                                      |                                     |                                     |                                     |                                      |                                      |  |
| Sequence 6654, Ap<br>Sequence 11079, A<br>Sequence 6406, Ap<br>Sequence 25623, A<br>Sequence 2, Appl<br>Sequence 2, Appl  | Sequence 10198, A<br>Sequence 120, App  | Patent No. 5244657<br>Patent No. 5243945 | Sequence 12846, A<br>Patent No. 5223610                                  | Patent No. 5244657<br>Patent No. 5433945                       | Sequence 24, Appl<br>Sequence 24, Appl                                    | Sequence 24, Appl                    | Sequence 19233, Ap                     | Patent No. 5223610<br>Sequence 24307, A                          | Sequence 23778, A                        | Sequence 1323/, A<br>Sequence 17215, A  | Sequence 42251, A                       | Sequence 13934, A<br>Sequence 14, Appl | Sequence 14, Appl<br>Sequence 6. Appli                                   | Sequence 6, Appli                   | Sequence 818, App<br>Sequence 5479, Ap  | Sequence 5631, Ap                       | Sequence 20, Appl                    | Sequence 26, Appl<br>Sequence 20, Appl                                       | Sequence 19408, A   | Sequence 12, Appl                     | sequence 17, Appi<br>Sequence 12551, A  | Sequence 96, Appl                    | Sequence 1, Appli                    | Sequence 7795, Ap                       | Sequence 8329, Ap                       | Sequence 15524, A                       | Sequence 4908, Ap                       | Sequence 7214, Ap                       | Sequence 113, APP<br>Sequence 6548, Ap | Sequence 1, Appli                   | Sequence 284, App                      | Sequence 1490, Ap                       | Sequence 1492, Ap<br>Sequence 26246. A                            | Sequence 4795, Ap                       | Sequence 2, Appli                   | Sequence 143/0, A                    | Sequence 2, Appli                   | Sequence 8, Appli                   | Sequence 3, Appli                   | Sequence 8, Appli                    | Sequence 3, Appli                    | Sequence 3, Appii<br>Sequence 2, Appli |
| Sequence<br>Sequence<br>Sequence<br>Sequence  | Sequence  | Patent No<br>Patent No                   | Sequence<br>Patent No  | Patent No  | Sequence  | Sequence                             | Sequence                               | Patent M<br>Seguence   | Sequence                                 | Sequence                                | Sequence                                | Sequence                               | Sequence   | Sequence                            | Sequence  | Sequence                                | Sednence                             | Sequence   | Sequence  | Sequence                              | Sequence  | Sequence                             | Sequence                             | Sequence                                | Sequence                                | Sequence                                | Sequence                                | Sequence                                | Sequence                               | Sequence                            | Sequence                               | Sequenc                                 | Seguenc   | Sequence                                | Sequence                            | Sequenc                              | Sequence                            | Seguence                            | Sequenci                            | Seguence                             | Sequenc                              | Sequence                               |
| 2 US-09-543-681A-6654 Sequence<br>2 US-09-902-540-11079 Sequence<br>2 US-09-681A-6406 Sequence<br>2 US-09-071-035-120 Sequence<br>2 US-09-291A-26823 Sequence<br>2 US-09-990-337-2 Sequence   | 2 US-09-902-540-10198 Sequence<br>2 US-10-206-576-120 Sequence<br>2 US-10-206-576-120 Sequence  | 6 5244657-6 5244657-6 5433945-6 Patent N | 2 US-09-902-540-12846 Sequence<br>6 5223610-6 Patent N                   | 6 5244657-9 Patent No 6 5433945-9 Patent No                    | 2 US-09-457-568-24 Sequence 2 US-09-457-646-24 Sequence                   | 2 US-09-516-065-24 Sequence          | 2 US-10-104-047-3435 Sequence          | 6 5223610-9<br>2 US-09-248-796A-24307 Sequence                   | 2 US-09-252-991A-23778 Sequence          | 2 US-09-252-991A-17215 Sequence         | 2 US-09-270-767-42251 Sequence          | 2 US-09-354-129-14 Sequence            | 2 US-09-504-357-14 Sequence<br>2 US-09-167-434-6 Sequence                | 2 US-08-853-755-6 Sequence          | 2 US-09-198-452A-818 Sequence<br>2 US-09-134-000C-5479 Sequence                   | 2 US-09-134-001C-5631 Sequence          | 2 US-09-457-646-20 Sequence          | 2 US-08-902-572-26 Sequence<br>2 US-09-516-065-20 Sequence                   | 2 US-09-252-991A-19408 Sequence<br>2 US-09-252-991A-19656 Sequence                | 1 US-08-114-555A-12 Sequence          | 2 US-08-559-39/A-1/<br>2 US-09-489-039A-12551 Sequence                            | 2 US-10-012-819-96 Sequence          | 2 US-09-353-242A-1 Sequence          | 2 US-09-489-039A-7795 Sequence          | 2 US-09-543-681A-8329 Sequence          | 2 US-09-902-540-15524 Sequence          | 2 US-09-107-532A-4908 Sequence          | 2 US-09-489-039A-7214 Sequence          | 2 US-09-949-016-6548 Sequence          | 2 US-09-987-107-1 Sequence          | 2 US-10-020-445A-284 Sequence          | 2 US-09-605-703B-1490 Sequence          | 2 US-09-605-703B-1492 Sequence<br>2 US-09-252-991A-26246 Sequence | 2 US-09-134-000C-4795 Sequence          | 2 US-09-987-107-2 Sequence          | 2 US-10-104-047-3660 Sequence        | 1 US-08-463-911-2 Sequence          | 2 US-09-140-804-8 Sequence          | 2 US-09-118-408-3 Sequence          | 2 US-09-686-838B-8 Sequence          | 2 US-09-911-176B-3 Sequence          | 2 US-09-776-976-2 Sequence             |
| 5 11.6 224 2 US-09-543-681A-6654 Sequence 6654, April 1.6 224 2 US-09-902-540-11079 Sequence 1079, 3 11.6 225 2 US-09-543-681A-6406 Sequence 6406, April 1.6 226 2 US-09-71-035-120 Sequence 120, April 226 2 US-09-252-991A-26823 Sequence 26823, 4 25 2 US-09-990-337-2 Sequence 26823, 4 April 226 2 US-09-990-337-2 | 11.6 226 2 US-09-902-540-10198 Sequence 11.6 226 2 US-09-902-576-120 Sequence 11.6 226 2 US-08-6776-120 Sequence 11.6 200-200-200-200-200-200-200-200-200-200 | 11.6 226 6 5433945-6 Patent N            | 11.6 227 2 US-09-902-540-12846 Sequence<br>11.6 227 6 5223610-6 Patent N | 11.6 227 6 5244657-9 Patent N<br>11.6 227 6 5433945-9 Patent N | 11.6 228 2 US-09-457-568-24 Sequence 11.6 228 2 US-09-457-646-24 Sequence | 11.6 228 2 US-09-516-065-24 Sequence | 11.6 228 2 US-10-104-047-3435 Sequence | 11.6 228 6 5223610-9<br>11.6 229 2 US-09-248-796A-24307 Sequence | 11.6 230 2 US-09-252-991A-23778 Sequence | 11.6 233 2 US-09-502-540-1525/ Sequence | 11.6 233 2 US-09-270-767-42251 Sequence | 11.6 234 2 US-09-354-129-14 Sequence   | 11.6 234 2 US-09-504-357-14 Sequence 11.6 235 2 US-09-167-434-6 Sequence | 11.6 235 2 US-08-853-755-6 Sequence | 11.6 235 2 US-09-198-452A-818 Sequence<br>11.6 235 2 US-09-134-000C-5479 Sequence | 11.6 237 2 US-09-134-001C-5631 Sequence | 11.6 237 2 US-09-457-646-20 Sequence | 11.6 237 2 US-08-902-572-26 Sequence<br>11.6 237 2 US-09-516-065-20 Sequence | 11.6 237 2 US-09-252-991A-19408 Sequence 11.6 237 2 US-09-252-991A-19656 Sequence | 11.6 239 1 US-08-114-555A-12 Sequence | 11.5 239 2 US-U8-559-39/A-1/ Sequence<br>11.6 239 2 US-09-489-039A-12551 Sequence | 11.6 239 2 US-10-012-819-96 Sequence | 11.6 241 2 US-09-353-242A-1 Sequence | 11.6 241 2 US-09-489-039A-7795 Sequence | 11.6 242 2 US-09-543-681A-8329 Sequence | 11.6 242 2 US-09-902-540-15524 Sequence | 11.6 243 2 US-09-107-532A-4908 Sequence | 11.6 243 2 US-09-489-039A-7214 Sequence | 11.6 243 2 US-09-949-016-6548 Sequence | 11.6 243 2 US-09-987-107-1 Sequence | 11.6 243 2 US-10-020-445A-284 Sequence | 11.6 243 2 US-09-605-703B-1490 Sequence | 11.6 244 2 US-09-252-991A-26246 Sequence                          | 11.6 244 2 US-09-134-000C-4795 Sequence | 11.6 244 2 US-09-987-107-2 Sequence | 11.6 246 2 US-104-340-149/8 Sequence | 11.6 247 1 US-08-463-911-2 Sequence | 11.6 247 2 US-09-140-804-8 Sequence | 11.6 247 2 US-09-118-408-3 Sequence | 11.6 247 2 US-09-686-838B-8 Sequence | 11.6 247 2 US-09-911-176B-3 Sequence | 11.6 247 2 US-09-716-976-2 Sequence    |

2, Appli 43923, A 2, Appli 8, Appli

us-10-624-218-3.oligo.rai

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TYPE: PRT
; ORGANISM: Drosophila melanogaster
US-09-270-767-42120
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ORGANISM: Pseudomonas aeruginosa
                TYPE: PRT
ORGANISM: Klebsiella pneumoniae
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Best Local Similarity 100...
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Best Local Similarity 100.
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Best Local Similarity 100.
Matches 7; Conservative
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91 RQRVQRL 97
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LENGTH: 323
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APPLICANT: Gary Breton et. al
APPLICANT: Gary Breton et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: NUCLEIC FOR DIAGNOSTICS AND THERAPEUTICS
TITLE REFERENCE: 2709.2004001
FILE REFERENCE: 2709.2004001
CURRENT APPLICATION NUMBER: US, 69/489, 039A
CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US, 60/117,747
PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
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6112, Ap
10229, A
31, Appl
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12, Appl
9, Appli
10, Appl
11, Appl
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APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
TITLE OF INVENTION: Nucleic 37326-094
CURRENT APPLICATION NUMBER: 1379-03-17
NUMBER OF SEQ ID NOS: 62217
NOWHER OF SEQ ID NOS: 62217
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 42478
LENGTH: 296
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Sequence 8, Appli
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US-09-253-682-16
US-09-457-0408-39
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US-09-252-991A-26366
US-09-987-107-10
US-09-987-107-10
US-09-987-107-11
US-09-3087-107-11
US-09-3087-107-11
US-09-3087-1029
US-09-3087-1029
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US-10-152-706-31
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US-08-55-378A-88
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100.0%; Pred. No. 1.4;
tive 0; Mismatches
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US-09-270-767-42478
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; Sequence 11444, Application US/09489039A
; Patent No. 6610836
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; Sequence 42478, Application US/09270767
; Patent No. 6703491
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ORGANISM: Drosophila melanogaster
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Best Local Similarity 100.vv
--hen 8; Conservative
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## Sequence 31594, Application US/09252991A

## SPELCANT: Marc J. Rubenfield et al.

## TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS

## TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

## TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

## CURRENT APPLICATION NUMBER: US/09/252,991A

## CURRENT FILING DATE: 1999-02-18

## PRIOR FILING DATE: 1998-02-18

## PRIOR FILING DATE: 1998-07-27

## NUMBER OF SEQ ID NOS: 33142

## LENGTH: 217
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Patent No. 6703491

GENERAL INFORMATION:
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
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                                              Indels
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16.3%; Score 7; DB 2;
100.0%; Pred. No. 10;
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100.0%; Pred. No. 16;
tive 0; Mismatches
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100.0%; Pred. No. 11;
tive 0; Mismatches
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Sequence 2486 Application US/0925291A

TITLE OF INVENTION: UNCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS

TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

CURRENT APPLICATION NUMBER: US 60/094,190

PRIOR FILING DATE: 1998-02-18

NUMBER OF SEQ ID NOS: 33142

SEQ ID NO 24868

LENGTH: 485
                                                                                                                                                                                                                                                                                                                                                                                         US-09-949-016-8999

Sequence 8999, Application US/09949016

Patent No. 6812339

HINTERMATION:
APPLICANT: VEWTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF;
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF;
CURRENT PILLING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-00-06

PRIOR FILING DATE: 2000-00-06

PRIOR FILING DATE: 2000-00-06

PRIOR FILING DATE: 2000-00-06

PRIOR FILING DATE: 2000-00-09

SOUPHARE: FASTESEQ for Windows Version 4.0
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                                                        Length 477;
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                                                        Query Match 16.3%; Score 7; DB 2; Best Local Similarity 100.0%; Pred. No. 24; Matches 7; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16.3%; Score 7; DB 2
100.0%; Pred. No. 24;
Live 0; Mismatches
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Matches 7; Conservative
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                                                                                                                                                                                                                                                                     197 QRVQRLR 203
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                                                                                                                                                                                                  24 QRVQRLR 30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 8
US-09-252-991A-24868
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US-09-252-991A-24868
US-09-252-991A-20516
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 8999
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APPLICANT: Marc J. Rubenfield et al.
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REPERBUCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT PILING DATE: 1999-02-18
PRIOR PLILING DATE: 1999-02-18
PRIOR PLILING DATE: 1998-01-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 2016
       Sequence 35, Application US/08311731A
| Sequence 35, Application US/08311731A
| Patent No. 6583266
| GENERAL INFORMATION:
| APPLICANT: SMITH, DOUGLAS
| TITLE OF INVENTION: NELATING TO MYCOBACTERIUM TUBERCULOSIS AND LAPRAE FOR TITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS
| TITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS NUMBER OF SEQUENCES 411
| CORRESPONDENCE ADDRESS: 411
| ADDRESSEE: WOLF, GRREENFIELD & SACKS, P.C. STREET: 600 ATLANTIC AVENUE CITY: BOSTON |
| STREET: MASSACHUSETTS |
| COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMFUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/311,731A
FILING DATE:
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: GATES, EDWARD R.
REFERENCES/DOCKET NUMBER: C0044/7125
TELECOMMUNICATION INFORMATION:
TELEFEHONE: 617/720-3500
TELEFEHORE: 617/720-3500
TELEFEHORE: 617/720-341
INFORMATION FOR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
LENGTH: 341 amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: ACID NO: ACID NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: protein HYPOTHETICAL: YES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        95 IEAAFLE 101
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US-09-252-991A-20516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-311-731A-35
   US-08-311-731A-35
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RESULT 11
US-08-630-349-5
Sequence 5, Application US/08630349
Patent No. 573908
Patent No. 573908
CENERAL INFORMATION:
TILLE OF INVENTION:
CORRESPONDENCE ADDRESS:
ADDRESSE:
ADDRESSE:
ADDRESSE:
ADDRESSE:
ADDRESSE:
ADDRESSE:
COUNTRY:
COUNTRY:
COUNTRY:
COUNTRY:
COUNTRY:
COUNTRY:
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 144 mb
COMPUTER: IBM COMPATIBLE
COMPUTER: IBM COMPATIBLE
COMPUTER: SYSTEM:
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FILING DATE:
PUBLICATION DATE:
RELEVANT RESIDUES IN SEQ ID NO:
US-08-285-440-5
    LENGTH: 532 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY:
LOCATION:
LDCATION:
LDENTIFICATION METHOD:
OTHER INFORMATION:
AUTHORS:
                                                                                                                                                       MOLECULE TYPE: peptide
HYPOTHETICAL:
ANTI-SENSE:
FRACHENT TYPE:
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 100.
Matches 7; Conservative
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STRAIN:
INDIVIDUAL ISOLATE:
DEVELOPMENTAL STAGE:
HAPLOTYPE:
TISSUE TYPE:
CELL TYPE:
CELL LINE:
CELL LINE:
IMMEDIATE SOURCE:
LIBRARY:
CLONE:
POSITION IN GENOME:
CHROMOSOME/SEGMENT:
MAP POSITION:
                                                                                   single
                                         TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
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                                                                                                    US-08-942-012B-24

US-08-942-012B-24

J Sequence 24, Application US/08942012B

Patent No. 6235278

GENERAL INFORMATION:

APPLICANT: Miller, Lois K.

APPLICANT: Dierks, Peter

APPLICANT: Lu, Albert

APPLICANT: Black, Bruce

TITLE OF INVENTION: Blological Insect Control Agents Expressing

TITLE OF INVENTION: Insect-Specific Toxin Genes, Methods and Compositions

FILE REFERENCE: 28-96a

CURRENT FILING DATE: 1997-10-01

PRIOR APPLICATION NUMBER: US/08/942,012B

CURRENT FILING DATE: 2000-10-01

PRIOR FILING DATE: 2000-10-01

NUMBER OF SEQ ID NOS: 33

SOFTWARE: PatentIn Ver. 2.0

SEQ ID NO 24

LENGTH: 515

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; ORGANISM: Helicoverpa zea nuclear polyhedrosis virus
US-08-942-012B-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 5, Application US/08285440
Patent No. 553237
GENERAL INFORMATION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSEE: Wenderoth, Lind & Ponack
STREET: 805 Fifteenth Street, N.W., #700
CITY: Weashington
STATE: D.C.
COUNTRY: U.S.A.
ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Discette, 5.25 inch, 500 kb
COMPUTER: IBM Compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: DisplayMrite
CURRENT APPLICATION NUMBER: US/08/285,440
FILING DATE:
CREATION NUMBER: US/08/285,440
FILING DATE:
CREATION NUMBER: US/08/285,440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16.3%; Score 7; DB 2;
100.0%; Pred. No. 25;
iive 0; Mismatches
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/858,947
FILING DATE: March 27, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Warren M. Cheek, Jr.
REGISTRATION NUMBER: 33,367
REFERENCE/DOCKET NUMBER:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX:
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 100.
Matches 7; Conservative
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428 EIEAAFL 434
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Length 532; 0; Indels

0; Gaps

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16.3%; Score 7; DB 2; Length 541; 100.0%; Pred. No. 27; tive 0; Mismatches 0; Indels
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Fatent No. 5532337
GENERAL INFORMATION:
APPLICANT: Ken'ichiro HAYASHI et al.
TITLE OF INVENTION: POLYBEPTIDE
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSEE: Wenderoth, lind & Ponack
STREET: Washington
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 500 kb
COMPUTER: IBM Compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: DisplayWrite
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/285,440
FILING DATE:
                FILE KEFEKELS CLOUD WINDER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-3
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-09
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-10-03
SOFTWARE: PRESEQ FOR WINDER: 6/231,498
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/858,94
FILING DATE: March 27,1992
ATTORNEY/AGENT INFORMATION:
NAME: WARTEN M. CHECK, JT.
REFERENCE/DOCKET NUMBER: 33,367
REFERENCE/DOCKET NUMBER:
TELECPHONE: 202-371-8850
TELECPHONE: 202-371-8850
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SEQUENCE CHARACTERISTICS:
LENGTH: 558 amino acids
TYPE: amino acid
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Best Local Similarity 10v..
---a 7; Conservative
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ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                    ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                            US-09-949-016-7217
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Patent No. 681239

GENERAL INFORMATION:

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF;

FILE REFERENCE: CLOOL337

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT PILING DATE: 2000-04-14

PRIOR FILING DATE: 2000-10-20

PRIOR FILING DATE: 2000-10-03

SECOFTWARE: PSECIENT NUMBER: 60/231,498

NUMBER OF SEC ID NOS: 207012

SECOFTWARE: PSECSE OF Windows Version 4.0

SECOFTWARE: PSECSE OF Windows Version 4.0
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Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
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100.0%; Pred. No. 26;
ive 0; Mismatches
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100.0%; Pred. No. 26;
Itive 0; Mismatches
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/630,349
FILING DATE: APALI 10, 1996
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/285,440
FILING DATE: August 4, 1994
PRIOR APPLICATION NUMBER: 07/858,947
FILING DATE: MARCH 27, 1992
ATTORNEY AGENT INFORMATION:
NAME: WARTEN M. Cheek, Jr.
REGISTRATION NUMBER: 33,367
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 532 amino acide
                                                                                                                                                                                                                                                                                                                                                               : 532 amino acids
amino acid
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Best Local Similarity 100.
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Matches 7; Conservative
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; MOLECULE TYPE: peptide
US-08-630-349-5
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; ORGANISM: Human
US-09-949-016-6479
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US-09-949-016-6479
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Query Match 16.3%; Score 7; DB 1; Length 558; Best Local Similarity 100.0%; Pred. No. 27; Matches 7; Conservative 0; Mismatches 0; Indels
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               NAME: Warren M. Cheek, Jr.
REGISTRATION NUMBER: 33,367
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 558 amino acids
TYPE: amino acid
STRANDENESS: single
FOPOLOGY: linear
NOLECTLE TYPE: peptide
US-08-630-349-6
ATTORNEY/AGENT INFORMATION:
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87 EAAFLER 93
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Fatent No. 573908
GENERAL INFORMATION:
APPLICANT: Ken'ichiro HAYASHI et al.
TITLE OF INVENTION:
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSEE: Wenderoth, Lind & Ponack
STREET: 805 Fifteenth Street, N.W., #700
CITY: Washington
STREET: B.C.
COUNTRY: U.S.A.
ZIP: 20005
COMPUTER EADDABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 144 mb
COMPUTER IBM Compatible
OPERATING SYSTEM: MS-DOS
SOFTWARE: Wordperfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/630,349
FILING DATE: April 10, 1996
CLASSIFTATION ATA:
APPLICATION NUMBER: 08/285,440
FILING DATE: MAJURE 4, 1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/858,947
FILING DATE: MAJURE: 07/858,947
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DOCUMENT NUMBER:
FILING DATE:
PUBLICATION DATE:
RELEVANT RESIDUES IN SEQ ID NO:
US-08-285-440-6
                                                                                                                                                                                  CLOME:
POSITION IN GENOME:
CHROMOSOME/SEGMENT:
MAP POSITION:
UNITS:
FEATURE:
LOCATION:
LOCATION:
IDENTIFICATION METHOD:
OTHER INFORMATION:
PUBLICATION INFORMATION:
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Best Local Similarity 100.
Matches 7; Conservative
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DBVELOPMENTAL STAGE:
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CELL TYPE:
CELL LINE:
ORGANELLE:
IMMEDIATE SOURCE:
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|87 EAAFLER 93
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VOLUME:
ISSUE:
PAGES:
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Sequence 2989, A
Sequence 15722, A
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Sequence 17, Appl
Sequence 10, Appl
Sequence 10, Appl
Sequence 20760, A
Sequence 20760, A
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5, Appli
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                 Sequence 68,
Sequence 49,
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US-09-159-469-61
     11016, A
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/cgn2_6/ptcdata/1/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptcdata/1/iaa/RE_COMB.pep:*
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                 GenCore version 5.1.6
(c) 1993 - 2005 Compugen Ltd.
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US-09-570-337B-2
US-09-570-337B-3
US-09-570-337B-3
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US-09-570-337B-4
US-09-973-278-372
US-09-973-278-372
US-09-973-278-372
US-09-973-278-372
US-09-543-6B1A-4424
US-09-543-6B1A-4424
US-09-543-6B1A-4345
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US-09-902-540-13147
US-09-903-750-2
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Gapop 60.0 , Gapext 60.0
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seq length: 200000000
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| 36443, A                        | Appli   | Appli<br>Appli                      | Appli                      | Appli<br>Appli  | Appl Appl  |                            | a, App                         |                             | l, App                      | Appli                                  | Appli                     | Appli                     | Appii<br>Appi  | Appl                       | Appl                       | Appl                       | Appl                       | Appl                       | App1                                    | Appl                       | Appl                       | Appl                                | Appl                       | Appl                       | Appl                                  | Appli<br>Appli              | Appl                       | App1                                   | Appl                        | App1                       | App1                       | Appl                       | Appl                       | App1  | Appl                       | Appl                         | Appl                               | , Appl                       | Appl                         | Appl                                   | Appl                         | Appl                         | Appl                                   | App1                       | Appl                         | Appl                         | App1                         | Appl                         | App1                          | Appli                       | Appl<br>Appl   |
|---------------------------------|---|-------------------------------------|----------------------------|---|--|----------------------------|--------------------------------|-----------------------------|-----------------------------|--|---------------------------|---------------------------|--|----------------------------|----------------------------|----------------------------|----------------------------|----------------------------|---|----------------------------|----------------------------|-------------------------------------|----------------------------|----------------------------|---------------------------------------|-----------------------------|----------------------------|--|-----------------------------|----------------------------|----------------------------|----------------------------|----------------------------|---|----------------------------|------------------------------|------------------------------------|------------------------------|------------------------------|--|------------------------------|------------------------------|--|----------------------------|------------------------------|------------------------------|------------------------------|------------------------------|-------------------------------|-----------------------------|--|
| Φ (                             | equence 510<br>equence 7,   |                                     | 0                          | 4 (4  | Sequence 25,<br>Sequence 24.                             | 52                         | Sequence 178<br>Sequence 80.   | ·                           | Sequence 141                | Sequence 3,                            | Sequence 3,               | Sequence 7,               | Sequence 7,<br>Sequence 38.                              | 7                          | Sequence 24,               | Sequence 25.               | N                          | 0                          | Sequence 96,                            | Sequence 24,               | Sequence 26,               | Sequence 27,                        | 1 (1)                      | Н.                         | Sequence 96,                          | nω                          | σ,                         | Sequence 13,                           | 0                           | ~                          | Sequence 24,               | 9 60                       | ~                          | Sequence 29,  | N                          | o c                          | equence 27,                        | 7                            | equence 29,                  | Sequence 24.                           | 7                            | ~ ~                          | sequence 28,<br>Seguence 29,           | 0                          | 0 0                          | equence 27,                  | 6                            | equence 29,                  | 9 (0)                         | 9                           | equence 23,<br>equence 24.                                   |
| 11.6 29 2 US-09-270-767-36443 S | 2 US-09-2/0-/6/-51660 S<br>1 US-08-287-537B-7 S<br>1 TG-06-207-537B-9 | 11.6 30 4 PCT-US94-10936-7          | 11.6 31 2 US-08-390-353A-4 | 11.6 31 2 US-09-101-059-4<br>11.6 31 2 US-09-359-921-24 | 11.6 31 2 US-09-359-921-25<br>11.6 31 2 US-09-360-344-24 | 11.6 31 2 US-09-360-344-25 | 11.6 31 2 US-09-9/3-2/8-1/8    | 11.6 32 2 US-09-227-357-181 | 11.6 32 2 US-09-079-030-141 | 11.6 32 2 US-09-491-6145-3             | 11.6 32 2 US-09-603-885-3 | 11.6 33 1 US-08-430-633-7 | 11.6 33 1 US-08-936-854-7<br>11.6 33 1 US-08-690-011A-38 | 11.6 33 2 US-09-058-459-23 | 11.6 33 2 US-09-058-459-24 | 11.6 33 2 US-03-036-439-26 | 11.6 33 2 US-09-058-459-28 | 11.6 33 2 US-09-058-459-29 | 11.6 33 Z US-U9-U8Z-Z79B-96             | 11.6 33 2 US-09-127-926-23 | 11.6 33 2 US-09-127-926-26 | 11.6 33 2 US-09-127-926-27          | 11.6 33 2 US-09-127-926-29 | 11.6 33 2 US-09-425-585-13 | 11.6 33 2 US-09-315-304B-96           | 11.6 33 2 US-09-293-495F-30 | 11.6 33 2 US-09-834-784-96 | 11.6 33 2 US-09-953-321-13             | 11.6 33 2 US-09-350-641C-96 | 11.6 33 2 US-09-714-357-23 | 11.6 33 2 US-09-714-357-24 | 11.6 33 2 US-09-714-357-27 | 11.6 33 2 US-09-714-357-28 | 11.6 33 2 US-09-714-357-29<br>11 6 33 2 US-09-350-8412-96 | 11.6 33 2 US-10-057-552-23 | 11.6 33 2 US-10-057-552-24 S | 11.6 33 2 US-10-057-552-26 8       | 11.6 33 2 US-10-057-552-28 S | 11.6 33 2 US-10-057-552-29 S | 11.6 33 2 US-09-827-960-23             | 11.6 33 2 US-09-827-960-26 S | 11.6 33 2 US-09-827-960-27 S | 11.6 33 2 US-09-827-960-28             | 11.6 33 2 US-09-837-886-23 | 11.6 33 2 US-09-837-886-24 S | 11.6 33 2 US-09-837-886-20 S | 11.6 33 2 US-09-837-886-28 S | 11.6 33 2 US-09-837-886-29 S | 11.6 33 2 US-09-491-614B-28 S | 11.6 33 2 US-10-323-314-6 S | 11.6 33 2 US-09-812-034-23 S<br>11.6 33 2 US-09-812-034-24 S |
| 174                             | 176   | 178                                 | 180                        | 181   | 183  | 185                        | 186                            | 188                         | 189                         | 191                                    | 192                       | 193                       | 194  | 196                        | 197                        | 199                        | 200                        | 201                        | 202                                     | 204                        | 205                        | 206                                 | 208                        | 209                        | 210                                   | 212                         |                            | 214                                    | 216                         | 217                        | 218                        | 220                        | 221                        | 222   | 224                        | 225                          | 227                                | 228                          | 229                          | 231                                    | 232                          | 233                          | 235                                    | 236                        | 237                          | 239                          | 240                          | 241                          |                               |                             | 245<br>246   |
| equence (                       | a) a) a   | ednence                             | equence 4                  | equence 1040<br>equence 1353                            | equence  | equence 7849, A            | equence z1000<br>equence 5890. | equence 41                  | equence 51, A               | Sequence 2, Appii<br>Sequence 12643, A | equence 1                 | ednence                   | equence (  | equence (                  | equence 31                 | Segmence 105.              | equence 14                 | equence 14                 | equence 36                              | equence 40                 | equence 15                 | equence 4,                          | equence 4,                 | equence 4,                 | equence 57,                           | equence 5<br>equence 5      | equence 1,                 | equence 1,                             | equence 2                   | equence 2,                 | equence 12                 | ednence                    | equence 3                  | equence   | equence 8                  | equence 9,                   | equence 3,<br>equence 9,           | equence 11(                  | equence 84                   | equence 84                             | equence 84                   | equence 84                   | equence 84                             | equence 8                  | equence 84                   | equence 84                   | equence 1,                   | equence 84,                  | equence 1566,                 | equence 1573,               | Sequence 84, Appl<br>Sequence 84, Appl                       |
| US-09-693-542-61                | US-US-09-693-542-69<br>US-10-258-860-6<br>US-10-258-860-2             | US-08-408-519-2<br>PCT-US95-03552-2 | 0-258-860-4                | US-09-489-039A-10407<br>US-09-489-039A-13531            | 3-714-741-40<br>3-252-991A-32                            | 9-949-016-7849             | 1-252-391A-21<br>1-328-352-589 | 3-714-741-41                | US-08-853-659A-51           | US-09-902-540-12643                    | US-09-914-259-11          | US-09-710-279-2964        | US-09-036-98/A-6<br>US-09-370-700-6                      | US-09-603-207-6            | US-09-134-001C-3159        | US-08-955-138-105          | US-08-400-796-14           | US-08-248-839C-144         | US-US-69U-ULLA-36<br>17S-09-299-4955-36 | US-09-149-476-409          | US-08-530-792D-19          | US-08-383-753-4<br>IIS-08-586-772-4 | US-08-959-512-4            | US-09-512-983-4            | US-08-056-200-57<br>TIS-08-800-644-57 | US-09-438-185A-578          | US-08-690-011A-1           | US-09-299-495F-1<br>IIS-09-491-614R-29 | US-08-690-011A-2            | US-09-299-495F-2           | US-09-079-030-128          | US-08-690-011A-3           | US-08-690-011A-37          | US-09-299-495F-3<br>US-09-299-495F-37                     | US-09-403-861A-8           | US-07-684-965-9              | US-08-737-829-3<br>US-09-252-658-9 | US-09-902-540-11089          | US-08-486-099-84             | US-08-380-10/A-94<br>US-08-484-223B-84 | US-08-919-597-84             | US-08-475-668A-84            | US-08-483-531A-84<br>US-08-471-913A-84 | US-08-485-264A-84          | US-08-474-349A-84            | US-08-470-896-84             | US-09-914-259-1              | US-08-485-546A-84            | US-09-350-841A-1566           | US-09-350-841A-1573         | US-08-487-266A-84<br>US-08-484-741-84                        |
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| 11.0.   10.0   | 11   11   11   12   13   14   14   15   15   15   15   15   15   | -182<br>-182<br>-182<br>-183<br>-317  | -469<br>G-79<br>A-17   | A-22<br>A-19<br>-109  | -321  | -515<br>-304<br>R-26   | D-82  | A-18   | A-18<br>-421                            | -337<br>-490<br>-20  | -112<br>-178  | -178  | -113<br>A-20  | -684<br>A-25   | A-16<br>A-48  | -365  | D-12  | -5<br>4-32  | -210   | -210   | A-26  | -550  | 2-35  | B-10   | -106   | -106<br>B-10   | -317                                       | -467  | -595  | -155                                      | A-20  | -12   |
| 11.0   10.0      | 11   11   11   12   13   14   14   15   15   15   15   15   15   | -325<br>-325<br>-325<br>-035<br>-039  | -767<br>-236<br>-999<br>-991   | -211<br>-796<br>-210  | -767  | -767<br>-639<br>-543   | -518  | -991   | -976                                    | -767   | -035  | -011  | -540  | -352<br>-796<br>-809   | -991  | -767  | -518  | - 395<br>- 395<br>- 1995  | -325   | -542   | -991  | -767  | -000  | -700   | -537   | -879<br>-043   | -767                                       | -540  | -767  | -540                                      | -7147   | - 04  |
| 11.0.   10.0.2   10.0.2527.7557.75186   80000000   2556  | 11   11   11   12   13   14   14   15   15   15   15   15   15   | -736<br>-902<br>-902<br>-017<br>-270  | -270<br>-345<br>-513<br>-252   | -640<br>-248<br>-732  | -270  | -270<br>-774<br>-249   | -303  | -252   | -201                                    | -270   | -071<br>-148  | -621  | -902  | -328<br>-248<br>-010   | -252  | -270  | -303  | -101<br>-673<br>-252  | -461   | -012   | -252  | -270  | -134  | -956   | -476   | -475<br>-433   | -270                                       | -902  | -270  | -902                                      | -602  | 3   |
| 11.0   10.0   20.0      | 11.0.   10.0.   2.0.   10.0. |   |  |   |   |  | က်လွှင်   | တ်တွင  | ກ່ວ່                                    | က်တ်လ  | က်လုတ်  | က်တွင်  | တ်တ   | က်တွင်   | တ်လ   | က်တွ  | 8 8   | ກຸ່ ໝໍ ໝໍ   | ວ ຕໍ່ ຕໍ່  | က်လ်   | ຜ່ວ   | က်လွ  | တ်ပ   | ່ເວັດ  | ည်တွင်   | င်္ပ င်္ပ  | က်ပ  | ည်တွင်  | SS  | က်င                                       | ກ່ວຸດ   | S-0   |
| 11.0   10.0      | 11.0   10.2   2.0   2. |   |  |   |   | ט ט ט  | נננ   | , c, c   | ; ر; ر                                  | ין נין ני  | ין נין נ  | ניני  | כנ  | םםם  | ם כן נ  |   | ם ם   | _ ;; c  | , ,,   | 9 12 12  | ; ;; ;  | , ,   | , p =   | ) (C) (                                      | ים כן  | ם  |  | -   |   |   |   |   |
| 11.6   10.2   10.6   10.5      | 11.0   10.2   2.0   10.0   4 | 1111122   | 443  | 47  | 949   | 9 4 9  | 51  | 54.  | 54.4                                    | 54   | 555   | 55  | 57  | 57.  | 9 8 8   | 288   | 86.0  | 969   | 36.6   | 19.5   | 2 5   | 200   | 9 4 4   | . 69   | 5 5 5  |  | 99.  |   | 99.   | 80  | 900   | 70  |
| 11.6   11.2   11.0      | 11.0   10.2   2.05.0-246-2956-2335   Sequence 233.6, p.     11.1   10.2   2.05.0-246-2956-2335   Sequence 233.6, p.     11.1   10.2   10.5-246-2956-2335   Sequence 23.7, pp.     11.1   10.2   10.5-246-2957-2027   Sequence 23.7, pp.     11.1   10.2   10.5-246-2957-2027   Sequence 23.7, pp.     11.1   10.2   10.5-246-296-296-296-296-296-296-296-296-296-29  |   |  |   |   |  |   |  | -, c1 (                                 |  |   |   | ан  | e4 e4 e4   |   |   |   | -,  |  |  |   |   |   | ·  |  | <del>-</del>   |  | - H   |   |   | ·   | 7 [   |
| 11.6   10.2   10.0-10.750-8664-18   Sequence 33.96   April     | 11.0   10.2   10.0-0.248-778-788   Sequence 731, Ap   468     11.0   10.2   10.0-0.248-778-784   Sequence 731, Ap   468     11.0   10.4   10.0-0.248-778-642   Sequence 731, Ap   468     11.0   10.4   10.0-0.248-778-642   Sequence 731, Ap   470     11.0   10.4   10.0-0.248-778-743   Sequence 731, Ap   470     11.0   10.5   10.0-0.248-778-743   Sequence 731, Ap   470     11.0   10.0   10.0-0.248-778   Sequence 731, Ap   470     11.0   10.0   10.0-0.248-778   Sequence 731, Ap   470     11.0   10.0   10.0-0.248-778   Sequence 731, Ap   480     11.0   11.0   10.0-0.248-778   Sequence 731, Ap   500     11.0   11.0   10.0-0.248-778   Sequence 731, Ap   500     11.0   11.0   11.0   11.0   11.0     11.0   11.0   11.0   11.0   11.0     11.0   11.0   11.0   11.0   11.0     11.0   11.0   11.0   11.0   11.0     11.0   11.0   11.0   11.0   11.0     11.0   11.0   11.0   11.0   11.0     11.0   11.0   11.0   11.0   11.0     11.0   11.0   11.0   11.0     11.0   11.0   11.0   11.0     11.0    |   |  |   |   |  |   |  |   |  |   |   |   |  |   |   |   |   |  |  |   |   |   |  |  |  |  | ا با  | 4   | ; ;; ,                                    | i.i.  |   |
| 11.6   10.2   10.0-0.0.0.0.0.0.0.0.0.0.0.0.0.0.0.0.0.0   | 111   111   112   113  |   |  | <b></b>   |   |  |   |  |   |  |   |   |   |  |   |   |   |   |  |  |   |   |   |  |  |  |  |   |   |   |   | 010   |
| 11.6   10.2   10.9   10.9   24.9   24.8   25.8   24.9   25.8   24.9   25.8   24.9      | 11.6   10.02 2   10.02-266-7958A-22368   Sequence 275, Appl.   10.00 1   10.00-267-795A-22368   Sequence 275, Appl.   10.00 1   10.00-267-795A-2358   Sequence 275, Appl.   10.00 1   10.00-277-795-70-755- |   |  |   |   |  |   |  |   |  |   |   |   |  |   |   |   |   |  |  |   |   |   |  |  |  |  |   |   |   |   |   |
| 11.6   10.2   10.5      | 11.6   10.02 2   10.02-266-7958A-22368   Sequence 275, Appl.   10.00 1   10.00-267-795A-22368   Sequence 275, Appl.   10.00 1   10.00-267-795A-2358   Sequence 275, Appl.   10.00 1   10.00-277-795-70-755- | <b>८</b> ळ००स८≀   | <b>ህ 44 (U (</b> 0)  | V 80 60   | о <del>г</del>  | 0 m 4  | • 15 19   | <b>~ 8</b> 0   | י כי ע                                  | 40.  | . 4 ru  | 9.  | 86  | 010  | : W 4   | <b>د</b> د د د  | <b>7</b> 8 7  | y O -   | 10 "   | ) 4r (C  |   | · co o  | 10 -  | 401  | <b>መ</b> 4 ፡   | ഗയ   |  | 20 ON   | 0 +   | 101                                       | ካ <b>ላ</b> ተ ሀ  | nω  |
| 11.6   10.2   1 US-09-7548-7565-716   Sequential (1)   10.2   1 US-09-748-7565-716   Sequential (1)   10.2   10.90-9-12-990C-837   Sequential (1)   10.4   2 US-09-12-990C-837   Sequential (1)   10.5   1 US-09-73-200-2   Sequential (1)   1 US-09-200-2   Sequential (1)   1 US-09-200-2   Sequential (1)   1 US-09   | 11.6   10.2   10.9   24.0   296.7   23.3   6   644.6   11.6   10.2   10.9   24.0   296.7   23.7   12   64.2   11.6   10.4   2   10.9   29.7   27.7   27.6   24.2   296.7   23.7   29.6 | 444444  | 4444   | 4<br>7<br>7<br>7  | 48  | 4 4 4<br>8 8 8   | 4 4 8   | 4 4 8 6  | 4. 4.<br>0 0                            | 4 4 4  | 4 4   | 44  | 4 9   | 2000   | 000   | 200   | 20  | 512   | 22.5   | 2010   | 51.   | 12.   | 22.5  | 22.0   | 22.23  | 22 22  | 22   | 22  | 2 23  | 200                                       | 200   | 20.00   |
| 11.6   10.2   1 USO-9-750-75386   Sequential Control of the cont   | 11.6   10.2   10.9   24.0   296.7   23.3   6   644.6   11.6   10.2   10.9   24.0   296.7   23.7   12   64.2   11.6   10.4   2   10.9   29.7   27.7   27.6   24.2   296.7   23.7   29.6 | , Appl<br>386, A<br>37, Ap<br>, Appl<br>2, App  | 4, App<br>Appli<br>44, Ap<br>167, Ap   | Appl<br>Appl<br>Appl  | Appl<br>Appl  | 8, App<br>, Appl<br>Appl   | 21, A<br>8, Ap  | App  | 8, Ap                                   | , Ap   | Appli<br>Appli  | 6, A  | Appl<br>Appli   | <u> </u>   | ppli<br>Ap  | 11.5  | 44  | App<br>App  | Ap   | A B  | dd:   | 13,8  | ÷ <u>0</u> . 9  | D. eC  | 0. 0.  | Ap<br>Ap   | jū.  | , A<br>App  | Ap<br>np.]  | App                                       | App   | •   |
| 111.6 102 2 US-08-755 111.6 103 1 US-08-755 111.6 111.6 113 1 US-08-755 111.6 111. | 11   6   102   2   103   | 0   | 9  | e 25<br>27<br>27  | 422   | 6 6 6 30 5 4 5 5 5 4 5 5 5 5 5 5 5 5 5 5 5 5 5   | e 211   | e 129  | 46.                                     | 9 60 -   | 3,  | 242   | 18,   | 8,<br>125<br>229   |   | . · w   | w m   | 136   | 26.5   | 151  | 120   | , 0, 4  | 1138,   | 2475   | 120,<br>573,   | 9217,<br>9218,   | 13, 7                                      | 287,  | 5156<br>14.   | 163,                                      | 163,  | 8121  |
| 111.6 102 2 US-08-755 111.6 103 1 US-08-755 111.6 111.6 113 1 US-08-755 111.6 111. | 11   6   102   2   103   | Sequence 16, Sequence 25, Sequence 27, Sequence 27, Sequence 27, Sequence 177, Sequence 177, Sequence 600   | Sequence 20<br>Sequence 20<br>Sequence 51<br>Sequence 40   | Sequence 29,<br>Sequence 25,<br>Sequence 27,  | 422   | Sequence 30<br>Sequence 24<br>Sequence 25  | Sequence 211<br>Sequence 447  | Sequence 129<br>Sequence 129   | 46.                                     | 9 60 -   | 3,  | 242   | 18,   | 8,<br>125<br>229   |   | . · w   | w m   | 136   | 26.5   | 151  | 120   | , 0, 4  | 1138,   | 2475   | 120,<br>573,   | 9217,<br>9218,   | 13, 7                                      | 287,  | 5156<br>14.   | 163,                                      | 163,  | 8121  |
| 11.6   10.2   10.5   0.8   0   | 11.6   10.2   2   10.5   10. | anties<br>anties<br>anties<br>anties  | antes<br>Sedne<br>Sedne  | Seque<br>Seque<br>Seque   | Sequence 24<br>Sequence 25  | Sequen<br>Sequen<br>Sequen   | Sequen  | Sequen   | Sequence 45                             | Sequence 48<br>Sequence 99   | Sequence 3,<br>Sequence 3,  | Sequence 24   | Sequence 18,  | Sequence 8,<br>Sequence 125<br>Sequence 229  | Sequence 1  | Sequence Sequence   | Sequence Sequence   | Sequence 2,<br>Sequence 136<br>Sequence 525   | Sequence 565   | Sequence 151   | Sequence 150  | Sequence 9,   | Sequence 1138,  | Sequence 2475                                | Sequence 120,<br>Sequence 573,                                       | Sequence 9217,<br>Sequence 9218,   | Sequence 13, 7                             | Sequence 13120<br>Sequence 287,   | Sequence 5156<br>Sequence 14.   | Sequence 163,                             | Sequence 553,   | Sequence 2509<br>Sequence 8121  |
|  |  | anties<br>anties<br>anties<br>anties  | antes<br>Sedne<br>Sedne  | Seque<br>Seque<br>Seque   | Sequence 24<br>Sequence 25  | Sequen<br>Sequen<br>Sequen   | Sequen  | Sequen   | Sequence 45                             | Sequence 48<br>Sequence 99   | Sequence 3,<br>Sequence 3,  | Sequence 24   | Sequence 18,  | Sequence 8,<br>Sequence 125<br>Sequence 229  | Sequence 1  | Sequence Sequence   | Sequence Sequence   | Sequence 2,<br>Sequence 136<br>Sequence 525   | Sequence 565   | Sequence 151   | Sequence 150  | Sequence 9,   | Sequence 1138,  | Sequence 2475                                | Sequence 120,<br>Sequence 573,                                       | Sequence 9217,<br>Sequence 9218,   | Sequence 13, 7                             | Sequence 13120<br>Sequence 287,   | Sequence 5156<br>Sequence 14.   | Sequence 163,                             | Sequence 553,   | Sequence 2509<br>Sequence 8121  |
|  |  | anties s  | antes<br>Sedne<br>Sedne  | Seque<br>Seque<br>Seque   | Sequence 24<br>Sequence 25  | Sequen<br>Sequen<br>Sequen   | Sequen  | Sequen   | Sequence 45                             | Sequence 48<br>Sequence 99   | Sequence 3,<br>Sequence 3,  | Sequence 24   | Sequence 18,  | Sequence 8,<br>Sequence 125<br>Sequence 229  | Sequence 1  | Sequence Sequence   | Sequence Sequence   | Sequence 2,<br>Sequence 136<br>Sequence 525   | Sequence 565   | Sequence 151   | Sequence 150  | Sequence 9,   | Sequence 1138,  | Sequence 2475                                | Sequence 120,<br>Sequence 573,                                       | Sequence 9217,<br>Sequence 9218,   | Sequence 13, 7                             | Sequence 13120<br>Sequence 287,   | Sequence 5156<br>Sequence 14.   | Sequence 163,                             | Sequence 553,   | Sequence 2509<br>Sequence 8121  |
|  |  | anties s  | antes<br>Sedne<br>Sedne  | Seque<br>Seque<br>Seque   | Sequence 24<br>Sequence 25  | Sequen<br>Sequen<br>Sequen   | Sequen  | Sequen   | Sequence 45                             | Sequence 48<br>Sequence 99   | Sequence 3,<br>Sequence 3,  | Sequence 24   | Sequence 18,  | Sequence 8,<br>Sequence 125<br>Sequence 229  | Sequence 1  | Sequence Sequence   | Sequence Sequence   | Sequence 2,<br>Sequence 136<br>Sequence 525   | Sequence 565   | Sequence 151   | Sequence 150  | Sequence 9,   | Sequence 1138,  | Sequence 2475                                | Sequence 120,<br>Sequence 573,                                       | Sequence 9217,<br>Sequence 9218,   | Sequence 13, 7                             | Sequence 13120<br>Sequence 287,   | Sequence 5156<br>Sequence 14.   | Sequence 163,                             | Sequence 553,   | Sequence 2509<br>Sequence 8121  |
|  |  | anties s  | antes<br>Sedne<br>Sedne  | Seque<br>Seque<br>Seque   | Sequence 24<br>Sequence 25  | Sequen<br>Sequen<br>Sequen   | Sequen  | Sequen   | Sequence 45                             | Sequence 48<br>Sequence 99   | Sequence 3,<br>Sequence 3,  | Sequence 24   | Sequence 18,  | Sequence 8,<br>Sequence 125<br>Sequence 229  | Sequence 1  | Sequence Sequence   | Sequence Sequence   | Sequence 2,<br>Sequence 136<br>Sequence 525   | Sequence 565   | Sequence 151   | Sequence 150  | Sequence 9,   | Sequence 1138,  | Sequence 2475                                | Sequence 120,<br>Sequence 573,                                       | Sequence 9217,<br>Sequence 9218,   | Sequence 13, 7                             | Sequence 13120<br>Sequence 287,   | Sequence 5156<br>Sequence 14.   | Sequence 163,                             | Sequence 553,   | Sequence 2509<br>Sequence 8121  |
| ਜ਼ਫ਼ਜ਼   |  | 1 US-08-750-856A-16 Seque<br>2 US-09-248-796A-25386 Seque<br>2 US-09-513-999C-7837 Seque<br>1 US-08-690-011A-25 Seque<br>2 US-09-227-357-172 Seque<br>2 US-09-227-357-172 Seque<br>2 US-09-270-757-60427 Seque  | 2.05-09-373-278-204 Seque<br>1 US-08-743-200-2 Seque<br>2 US-09-583-110-5144 Seque<br>2 US-09-107-433-4067 Seque   | 1 US-08-690-011A-29 Seque<br>2 US-09-299-495F-25 Seque<br>2 US-09-299-495F-27 Seque             | 1 US-08-710-749-24 Sequence 24<br>2 US-09-299-495F-29 Sequence 29 | 2 US-09-634-238-308 Sequen<br>2 US-09-147-875A-24 Sequen<br>2 US-09-147-875A-25 Sequen                         | 2 US-09-107-532A-4478 Sequen  | 2 US-08-545-809A-129 Sequen<br>2 US-09-515-697-129 Sequen<br>2 US-09-715-69-7129                         | 2 US-10-104-047-3918 Sequence 39        | 2 US-09-543-681A-482/ Sequence 48<br>2 US-09-902-540-9940 Sequence 99<br>1 US-08-08-1-13 Sequence 19                   | 2 US-09-367-206-18 Sequence 18  | 2 US-09-248-796A-24976 Sequence 24<br>2 US-09-806-437-2 Sequence 2,                 | 2 US-09-298-404-18 Sequence 18, 2 US-09-742-373-7 Sequence 7,                 | 2 US-09-742-373-8 Sequence 8,<br>2 US-09-489-039A-12512 Sequence 125<br>2 US-09-540-236-2295 Sequence 229                          | 2 US-09-806-437-1<br>2 US-09-513-999C-8075 Sequence                         | 2 US-10-394-058-7 Sequence 2 US-10-394-058-8 Sequence 8                 | 2 US-09-270-767-61783 Sequence 6 2 US-09-902-540-13274 Sequence 1 | 1 US-U8-64/1144.2 Sequence 2,<br>2 US-09-198-452A-136 Sequence 136<br>2 US-09-513-999C-5281 Semience 526            | 2 US-09-513-999C-5659 Sequence 565<br>2 US-09-32R-352-8093 Semience 805          | 2 US-09-248-796A-15103 Sequence 151<br>2 US-09-471-276-1460 Sequence 146         | 2 US-09-199-637A-150 Sequence 150                                       | 2 US-10-394-058-9 Sequence 9, 2 US-09-621-976-4973 Semi-part 497        | 2 US-09-732-210-1138 Sequence 1138, 2 US-09-732-813-4088 Sequence 1138, | 2 US-09-248-796A-24754 Sequence 2475         | 2 US-09-438-185A-120 Sequence 120, 2 US-10-198-053-573 Sequence 573, | 2 US-09-949-016-9217<br>2 US-09-949-016-9218 Sequence 9218,                                | 1 US-08-328-152A-13 Sequence 13, 2         | 2 US-09-502-540-13120 Sequence 1312(  | 2 US-09-107-433-5156 Sequence 5156<br>2 US-09-800-170-14 Sequence 14.                 | 2 US-09-904-615-163 Sequence 163,         | 2 US-US-10/1-312A-5359 Sequence 5555<br>2 US-10-054-988-163 Sequence 163, | 2 US-09-252-991A-26095 Sequence 2609<br>2 US-09-513-999C-8121 Sequence 8121         |
|  |  | 1 US-08-750-856A-16 Seque<br>2 US-09-248-796A-25386 Seque<br>2 US-09-513-999C-7837 Seque<br>1 US-08-690-011A-25 Seque<br>2 US-09-227-357-172 Seque<br>2 US-09-227-357-172 Seque<br>2 US-09-270-757-60427 Seque  | 2.05-09-373-278-204 Seque<br>1 US-08-743-200-2 Seque<br>2 US-09-583-110-5144 Seque<br>2 US-09-107-433-4067 Seque   | 1 US-08-690-011A-29 Seque<br>2 US-09-299-495F-25 Seque<br>2 US-09-299-495F-27 Seque             | 1 US-08-710-749-24 Sequence 24<br>2 US-09-299-495F-29 Sequence 29 | 2 US-09-634-238-308 Sequen<br>2 US-09-147-875A-24 Sequen<br>2 US-09-147-875A-25 Sequen                         | 2 US-09-107-532A-4478 Sequen  | 2 US-08-545-809A-129 Sequen<br>2 US-09-515-697-129 Sequen<br>2 US-09-715-69-7129                         | 2 US-10-104-047-3918 Sequence 39        | 2 US-09-543-681A-482/ Sequence 48<br>2 US-09-902-540-9940 Sequence 99<br>1 US-08-08-1-13 Sequence 19                   | 2 US-09-367-206-18 Sequence 18  | 2 US-09-248-796A-24976 Sequence 24<br>2 US-09-806-437-2 Sequence 2,                 | 2 US-09-298-404-18 Sequence 18, 2 US-09-742-373-7 Sequence 7,                 | 2 US-09-742-373-8 Sequence 8,<br>2 US-09-489-039A-12512 Sequence 125<br>2 US-09-540-236-2295 Sequence 229                          | 2 US-09-806-437-1<br>2 US-09-513-999C-8075 Sequence                         | 2 US-10-394-058-7 Sequence 2 US-10-394-058-8 Sequence 8                 | 2 US-09-270-767-61783 Sequence 6 2 US-09-902-540-13274 Sequence 1 | 1 US-U8-64/1144.2 Sequence 2,<br>2 US-09-198-452A-136 Sequence 136<br>2 US-09-513-999C-5281 Semience 526            | 2 US-09-513-999C-5659 Sequence 565<br>2 US-09-32R-352-8093 Semience 805          | 2 US-09-248-796A-15103 Sequence 151<br>2 US-09-471-276-1460 Sequence 146         | 2 US-09-199-637A-150 Sequence 150                                       | 2 US-10-394-058-9 Sequence 9, 2 US-09-621-976-4973 Semi-part 497        | 2 US-09-732-210-1138 Sequence 1138, 2 US-09-732-813-4088 Sequence 1138, | 2 US-09-248-796A-24754 Sequence 2475         | 2 US-09-438-185A-120 Sequence 120, 2 US-10-198-053-573 Sequence 573, | 2 US-09-949-016-9217<br>2 US-09-949-016-9218 Sequence 9218,                                | 1 US-08-328-152A-13 Sequence 13, 2         | 2 US-09-502-540-13120 Sequence 1312(  | 2 US-09-107-433-5156 Sequence 5156<br>2 US-09-800-170-14 Sequence 14.                 | 2 US-09-904-615-163 Sequence 163,         | 2 US-US-10/1-312A-5359 Sequence 5555<br>2 US-10-054-988-163 Sequence 163, | 2 US-09-252-991A-26095 Sequence 2609<br>2 US-09-513-999C-8121 Sequence 8121         |
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| Sequence 49088, A Sequence 27, Appl Sequence 27, Appl Sequence 27, Appl Sequence 1141, App Sequence 1833, A Sequence 1833, A Sequence 1, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 12, Appli Sequence 1497, App Sequence 16213, Appli Sequence 2844, Appli Sequence 2844, Appli Sequence 1141, Appli Sequence 1141, Appli Sequence 1141, Appli Sequence 10, Appli Sequence 22771, A Sequence 2771, A Sequence 22771, A Sequence 22771, A Sequence 22771, A Sequence 22771, A Sequence 1576, A Sequence 1576, A Sequence 2510, Appli Sequence 1576, Appli Sequence 1506, Appli Sequence 2510, Appli Sequence 26287, A   | ם ייייים ייייים ייייים ייייים יייייים יייייי   |
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| 281 2 US-09-538-092-221 Sequence 281 2 US-09-252-991A-19093 Sequence 282 2 US-09-252-991A-19093 Sequence 282 2 US-09-270-767-31445 Sequence 282 2 US-09-270-767-31999 Sequence 283 2 US-09-252-991A-2959 Sequence 283 2 US-09-252-991A-2959 Sequence 285 2 US-09-252-991A-2959 Sequence 286 2 US-09-262-991A-2959 Sequence 287 2 US-09-213-291A-2959 Sequence 288 2 US-09-213-291A-2959 Sequence 298 2 US-09-213-291A-2959 Sequence 299 2 US-09-213-340A-60 Sequence 299 2 US-09-223-341A-60 Sequence 299 2 US-09-223-341A-60 Sequence 299 2 US-09-223-341A-790 Sequence 299 2 US-09-223-341A-790 Sequence 299 2 US-09-223-341A-790 Sequence 299 2 US-09-223-341A-790 Sequence 290 2 US-09-223-341A-2329 Sequence 290 2 US-09-223-341A-2329 Sequence 290 2 US-09-223-341A-2329 Sequence 290 2 US   | 11.6 309 2 US-09-252-991A-23590 Sequence 23590<br>11.6 309 2 US-09-602-787A-294 Sequence 294,<br>11.6 309 2 US-09-248-796A-15871 Sequence 15873<br>11.6 309 2 US-09-902-540-15963 Sequence 15873<br>11.6 309 2 US-09-902-540-15963 Sequence 15963<br>11.6 309 2 US-10-206-576-108                            |
| 111.6 281 2 US-09-518-032-221 Sequence 2 111.6 281 2 US-09-528-0318-120 Sequence 2 111.6 281 2 US-09-270-767-12445 Sequence 2 111.6 282 2 US-09-270-767-12445 Sequence 2 111.6 282 2 US-09-270-767-14911 Sequence 2 111.6 282 2 US-09-543-618-74911 Sequence 2 111.6 282 2 US-09-270-767-14911 Sequence 2 111.6 282 2 US-09-543-618-74911 Sequence 2 111.6 282 2 US-09-543-618-74911 Sequence 2 111.6 282 2 US-09-543-618-74911 Sequence 2 111.6 285 2 US-09-543-618-74911 Sequence 2 111.6 286 2 US-09-543-618-74911 Sequence 2 111.6 286 2 US-09-743-74914 Sequence 2 US-   | 5 11.6 309 2 US-09-252-991A-23590 Sequence 23596<br>5 11.6 309 2 US-09-252-991A-23590 Sequence 234,<br>5 11.6 309 2 US-09-248-796A-15871 Sequence 294,<br>5 11.6 309 2 US-09-902-540-15963 Sequence 15873<br>5 11.6 309 2 US-09-902-540-15963 Sequence 15963<br>5 11.6 309 2 US-10-206-576-108 Sequence 108, |

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APPLICANT: VINSON, Charles R.

APPLICANT: WINSON, Charles R.

TITLE OF INVENTION: EXTENSION OF A PROTEIN-PROTEIN

TITLE OF INVENTION: INTERACTION SURFACE TO INACTIVATE THE FUNCTION OF A TITLE OF INVENTION: CELLULAR PROTEIN

NUMBER OF SEQUENCES: 60

CORRESPONDENCE ADDRESS:

ADDRESSEE: MORGAN & FINNEGAN, L.L.P.
                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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No. 0.044;
0; Indels
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SOFTWARE:

APPLICATION DATA:

RELIGNATION NUMBER:

FILING DATE:

APPLICATION NUMBER:

FILING DATE:

PRIOR APPLICATION NUMBER:

PLING DATE:

APPLICATION NUMBER:

PLING DATE:

APPLICATION NUMBER:

FILING DATE:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELEFAX:

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                                                                                                                                                                                                                         20.9%; Scor.
100.0%; Pred. No. ...
    CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 19335
SEQ ID NO 7606
LENGTH: 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: MORGAN & FINNEGAN, L.L.P. STREET: 345 Park Avenue CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-690-011A-52; Sequence 52, Application US/08690011A; Patent No. 5942433; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 10154-0053
ZOMPUTER READABLE PORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.
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                                                                                                                                                                        ; ORGANISM: Homo sapiens
US-09-621-976-7606
                                                                                                                                                                                                                                                                                                                                       21 ELEQEVORL 29
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Fatent No. 6812339

GENERAL INFORMATION:

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

CURRENT APPLICATION NUMBER: US/09/949,016

CURRENT APPLICATION NUMBER: 60/241,755

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR APPLICATION NUMBER: 60/231,498

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-00-08

NUMBER OF SEQ ID NOS: 207012

SOFTWARE: FastSEQ for Windows Version 4.0
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582, App
27226, A
14173, Ap
1050, Ap
1050, Ap
292, App
32, App
32, App
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23772, A
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                                                                                        US-09-902-540-14866
US-09-902-540-14866
US-09-605-703B-580
US-09-605-703B-582
US-09-543-681A-7707
US-09-543-681A-7707
US-09-543-681A-4835
US-09-543-681A-4835
US-09-949-016-1022
US-09-949-016-1022
US-09-902-540-10170
US-09-902-540-10170
US-09-902-540-10170
US-09-902-540-10170
US-09-913-881-32
US-09-913-881-32
US-09-912-881-32
US-09-913-881-32
US-09-913-881-32
US-09-913-881-32
US-09-913-881-32
US-09-913-881-32
US-09-913-881-32
US-09-913-881-32
US-09-913-881-32
US-09-913-913-913-6310
US-09-913-913-913-6310
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Patent No. 6639063
GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Jobert, S.
APPLICANT: Glordano, J.Y.
TITLE OF INVENTION: ESTS and Encoded Human Proteins.
FILE REFERENCE: GENSET.054PR2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-252-991A-23772
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ive 0; Mismatches
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Best Local Similarity 100.00
Conservative
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GENERAL INFORMATION:
APPLICANT: Buropean Molecular Biology Laboratory (EMBL)
APPLICANT: Buropean Molecular Helena
APPLICANT: Dominques, Helena
APPLICANT: Oregut, David
APPLICANT: Sebald, Walter
APPLICANT: Sebald, Walter
APPLICANT: Serveno, Luis
TITLE OF INVENTION: Ortokine Specific Receptors
FILE REFERENCE: 100564-00000
CURRENT APPLICATION NUMBER: US/09/570,337B
CURRENT FILING DATE: 2000-05-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: European Molecular Biology Laboratory (EMBL)
APPLICANT: European Molecular Biology Laboratory (EMBL)
APPLICANT: Cregut, David
APPLICANT: Sebald, Walter
APPLICANT: Sebald, Walter
APPLICANT: Serrano, Luis
APPLICANTION NUMBER: US/09/570,337B
CURRENT FILING DATE: 2000-05-12
NUMBER OF SEQ ID NOS: 9
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OTHER INFORMATION: Description of Artificial Sequence: synthetic peptide

NAME/KEY: misc feature

THER INFORMATION: Model 2, interleukin-4 mimetic peptide 4082 Kd micrometers

US-09-570-3378-2
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                                                                                                                                                                                                                      Length 425;
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100.0%; Pred. No. 21;
ive 0; Mismatches
                                                                                                                                                                                                                 Query Match 16.3%; Score 7; DB 2; Best Local Similarity 100.0%; Pred. No. 22; Matches 7; Conservative 0; Mismatches
  MBER: US 60/117,747
1999-01-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 3, Application US/09570337B Patent No. 6685932 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 2, Application US/09570337B ; Patent No. 6685932
PRIOR APPLICATION NUMBER: US 60/1
PRIOR FILING DAFF: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 11016
LENGTH: 425
TYPE: PRT
ORGANISM: Klebsiella pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 9
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
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Best Local Similarity 100.
Matches 6; Conservative
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                                                                                                                                                                       US-09-489-039A-11016
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US-09-570-337B-2
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Patent No. 6610836
GENERAL INFORMATION
GENERAL INFORMATION
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.2004001
CURRENT APPLICATION NUMBER: US/09/489,039A
CURRENT FILING DATE: 2000-01-27
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                                                                                                                                                                    APPLICANT: VINSON, Charles R.
APPLICANT: VINSON, DMILTY
TITLE OF INVENTION: EXTENSION OF A PROTEIN-PROTEIN
TITLE OF INVENTION: EXTENSION OF A PROTEIN
CELULAR PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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REGISTRATION NUMBER: 35,353
REFERENCE/DOCKET NUMBER: 2026-4199US2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 26-Apr-1999
PRIOR APPLICATION NUMBER: 60/001,654
APPLICATION NUMBER: 60/001,654
FILING DATE: 31-JUL-1995
APPLICATION NUMBER: 60/018,496
FILING DATE: 29-MAY-1996
APPLICATION NUMBER: 68/90,011
FILING DATE: 31-JULY-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 64
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORGAN & FINNEGAN, L.L.P.
STREET: 345 Park Avenue
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 2;
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Best Local Similarity 100.0%; Pred. No. 5;
Matches 7; Conservative 0; Mismatches
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TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 64:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 10154-0053
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: MS WORD 97
                                                                                                                     Sequence 64, Application US/09299495F
Patent No. 6361968
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)758-4800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 89 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          44 AELEQEV 50
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       44 ARLEQEV 50
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Gaps
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/note= "This amino acid is unknown."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 42;
                                                                                                                                     COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
FILING DATE: 19910829
CLASSIPICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION: 123 Human Secreted Proteins
TITLE OF INVENTION: 123 Human Secreted Proteins
FILE REFERENCE: PCD10P1
CURRENT APPLICATION NUMBER: US/09/227,357
CURRENT FILING DATE: 1999-01-08
EARLIER APPLICATION NUMBER: PCT/US98/13684
EARLIER FILING DATE: 1999-07-08
EARLIER FILING DATE: 1997-07-08
EARLIER APPLICATION NUMBER: 60/052,732
EARLIER APPLICATION NUMBER: 60/052,732
EARLIER FILING DATE: 1997-07-08
EARLIER APPLICATION NUMBER: 60/052,732
EARLIER FILING DATE: 1997-07-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                            ATTORNEY AGENT INFORMATION:
NAME: Sharkey, Richard G.
REGISTRATION NUMBER: 32,629
REFERENCE/DOCKET NUMBER: 150036.406C1
TELEPHONE: 206-622-4900
TELEPHONE: 206-622-4900
TELEPHONE: 206-682-6031
TELER: 3723836
INPORMATION FOR SEQ ID NO: 8:
SEGURNCE CHARACTERISTICS:
LENGTH: 42 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14.0%; Score 6; DB 1
100.0%; Pred. No. 27;
tive 0; Mismatches
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Patent No. 6342581
GENERAL INFORMATION:
                                                                                                                   E: Floppy disk
IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FRAGMENT TYPE: N-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 42 amino acids
TYPE: AMINO ACID
STRANDEDNESS: single
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Best Local Similarity 100..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                 ZIP: 98104
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION:
COTHER INFORMATION:
US-07-752-101A-8
                            Washington
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LOCATION: 28
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                                                 COUNTRY:
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APPLICANT: European Molecular Biology Laboratory (EMBL)
APPLICANT: Dominques, Helena
APPLICANT: Creatt, David
APPLICANT: Sebald, Walter
APPLICANT: Sebald, Walter
APPLICANT: Serrano, Luis
TITLE OF INVENTION: Design of Coiled-Coil Dimer Derived Antagonists of 4-Helix Bundle
TITLE OF INVENTION: Cytckine Specific Receptors
TITLE OF INVENTION: 100564-00000
CURRENT PAPLICATION NUMBER: 108/09/570,337B
CURRENT PLING DATE: 2000-05-12
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin version 3.0
SEQ ID NO 4
LENGTH: 31
                                                                                                                                 NAME/KEY: misc feature
OTHER INFORMATION: Description of Artificial Sequence: synthetic peptide
NAME/KEY: misc feature
OTHER INFORMATION: Model 2a, interleukin-4 mimetic peptide 2160 Kd micrometers
US-09-570-337B-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc feature;
CTHER INFORMATION: Description of Artificial Sequence: synthetic peptide
NAME/KEY: misc feature
NAME/KEY: Misc feature
US-09-570-337B-4
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Patent No. 5126857
GENERAL INFORMATION:
APPLICANT: Yamamoto, Fumi-ichiro
APPLICANT: White, Thayer
APPLICANT: Clausen, Henrik
TITLE OF INVENTION: ABO GENOTYPING
NUMBER OF SEQUENCES: 69
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
CORRESPENDENCE ADDRESS:
Seed and Berry
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                         14.0%; Score 6; DB 2;
100.0%; Pred. No. 21;
Live 0; Mismatches
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100.0%; Pred. No. 21;
iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 4, Application US/09570337B
Patent No. 6685932
SOFTWARE: PatentIn version 3.0
                                                               TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.
Matches 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6; Conservative
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Best Local Similarity
Matches 6; Conserva
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US-07-752-101A-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-570-337B-4
                     SEQ ID NO 3
LENGTH: 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
                                                                                                                FEATURE
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TITLE OF INVENTION: 123 Human Secreted Proteins
                                                                     CURRENT FILING DATE: 2001-10-10
REIOR PELICATION NUMBER: 09/227,357
REIOR PELICATION NUMBER: 09/227,357
REIOR PELICATION NUMBER: 09/227,357
REIOR PELICATION NUMBER: 09/227,357
REIOR PELICATION NUMBER: 09/051,926
REIOR APPLICATION NUMBER: 06/051,926
REIOR APPLICATION NUMBER: 06/051,926
REIOR APPLICATION NUMBER: 06/051,926
REIOR PELING DATE: 1997-07-08
REIOR RELING DATE: 1997-08-18
REIOR RELING 
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14.0%; Score 6; DB 2;
Best Local Similarity 100.0%; Pred. No. 42;
Matches 6; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EARLIER APPLICATION NUMBER: 60/055,954
EARLIER FILING DATE: 1997-08-18
EARLIER PELING DATE: 1997-08-18
EARLIER FILING DATE: 1997-09-12
SARLIER FILING DATE: 1997-09-12
SARLIER FILING DATE: 1997-09-12
SARLIER PILING DATE: 1997-09-12
SARLIER PILING DATE: 1997-09-12
SARLIER PILING DATE: 1997-09-12
SARLIER FILING DATE: 1997-09-12
SARLIER FILING DATE: 1997-09-12
SARLIER FILING DATE: 1997-09-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R APPLICATION NUMBER: 60/055,953
R FILING DATE: 1997-08-18
R FILING DATE: 1997-08-18
R APPLICATION NUMBER: 60/055,947
R FILING DATE: 1997-08-18
R APPLICATION NUMBER: 60/055,947
R APPLICATION NUMBER: 60/055,964
R PILING DATE: 1997-08-18
R APPLICATION NUMBER: 60/055,664
R FILING DATE: 1997-08-18
R APPLICATION NUMBER: 60/055,684
R FILING DATE: 1997-08-18
R APPLICATION NUMBER: 60/055,984
R FILING DATE: 1997-08-18
R PILING DATE: 1997-08-18
R PILING DATE: 1997-08-18
                             R PILING DATE: 1997-07-08
R APPLICATION NUMBER: 60/051,916
R PILING DATE: 1997-07-08
R PILING DATE: 1997-07-08
R PILING DATE: 1997-07-08
R APPLICATION NUMBER: 60/051,930
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R APPLICATION NUMBER: 60/052,733

R FILING DATE: 1997-07-08

R PILING DATE: 1997-07-08

R PILING DATE: 1997-07-08

R APPLICATION NUMBER: 60/051,919

R FILING DATE: 1997-07-08

R FILING DATE: 1997-07-08

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R FILING DATE: 1997-07-08
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R PAPLICATION NUMBER: 60/055,723
R PILING DATE: 1997-08-18
R APPLICATION NUMBER: 60/055,948
R PILING DATE: 1997-08-18
R PILING DATE: 1997-08-18
R PILING DATE: 1997-08-18
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US-09-973-278-372
; Sequence 372, Application US/09973278
; Patent No. 6924354
; GENERAL INFORMATION:
; APPLICANT: Fischer et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
CORGANISM: Homo sapiens
US-09-227-357-654
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                                                                                                     EARLIER
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Sequence 4345, Application US/09543681A

Patent No. 6605709
GENERAL INFORMATION:
APPLICANT: GARY BRETON
TITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS
CURRENT APPLICATION NUMBER: US/09/543,681A
CURRENT FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 60/128,706
PRIOR APPLICATION NUMBER: US 60/128,706
PRIOR FILING DATE: 1999-04-09
NUMBER OF SEQ ID NOS: 8344
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STREET: SALMON STREET
COUNTRY: USA
COUNTRY: USA
COUNTRY: USA
COUNTRY: USA
COMPUTER READABLE FORM:
WEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
WEDIUM TYPE: IBM PC compatible
COMPUTER: IBM PC compatible
COM
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APPLICANT: Parker, David B
APPLICANT: Process of the control of t
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100.0%; Pred. No. 54;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                     ; Sequence 10, Application US/08062472B
; Patent No. 5695954
; GENERAL INFORMATION:
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Best Local Similarity 100.
Matches 6; Conservative
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STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: peptide
            SOYETR 39
                                                                                  69 SOYETR 74
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Sequence 4424, Application US/09543681A
Sequence 4424, Application US/09543681A
Sequence 4424, Application US/09543681A
Setting Sequences Relating To Proteus Mirabil Title OF INVENTION: UCLEIC AND THERAPEUTICS
TITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: 1002-001
CURRENT APPLICATION NUMBER: US/09/543,681A
CURRENT FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 60/128,706
NUMBER OF SEQ ID NOS: 8344
SEQ ID NO 4424
LENGTH: 84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:

APPLICANT: Keith Weinstock et al
APPLICANT: Keith Weinstock et al
TITLE OF INVENTION: NUCLEIC ACID AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
CURRENT APPLICATION WUMBER: US/09/248,796A
CURRENT FILING DATE: 1999-02-12
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1998-08-13
NUMBER OF SEQ ID NOS: 28208
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100.0%; Pred. No. 43;
iive 0; Mismatches
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14.0%; Score 6; DB 2;
Best Local Similarity 100.0%; Pred. No. 53;
Matches 6; Conservative 0; Mismatches
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; Sequence 22740, Application US/09248796A
; Patent No. 6747137
                                                                                                                                                                                                                                                                                                                                                           14.0%; Scallarity 100.0%; Poc
Conservative 0;
PRIOR FILING DATE: 1997-09-12
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; ORGANISM: Proteus mirabilis
US-09-543-681A-4424
                            NUMBER OF SEQ ID NOS: 947
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 372
LENGTH: 66
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Best Local Similarity 100.
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                                                                                                                                                                                      TYPE: PRT
CRGANISM: Homo sapiens
US-09-973-278-372
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Best Local Similarity
Matches 6; Conserv
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; LENGTH: 91
; TYPE: PRT
; ORGANISM: Proteus mirabilis
US-09-543-681A-4345
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0; Gaps 0; Indels Ouery Match 14.0%; Score 6; DB 2; Length 91; Best Local Similarity 100.0%; Pred. No. 57; Matches 6; Conservative 0; Mismatches 0; Indels

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